

1/30

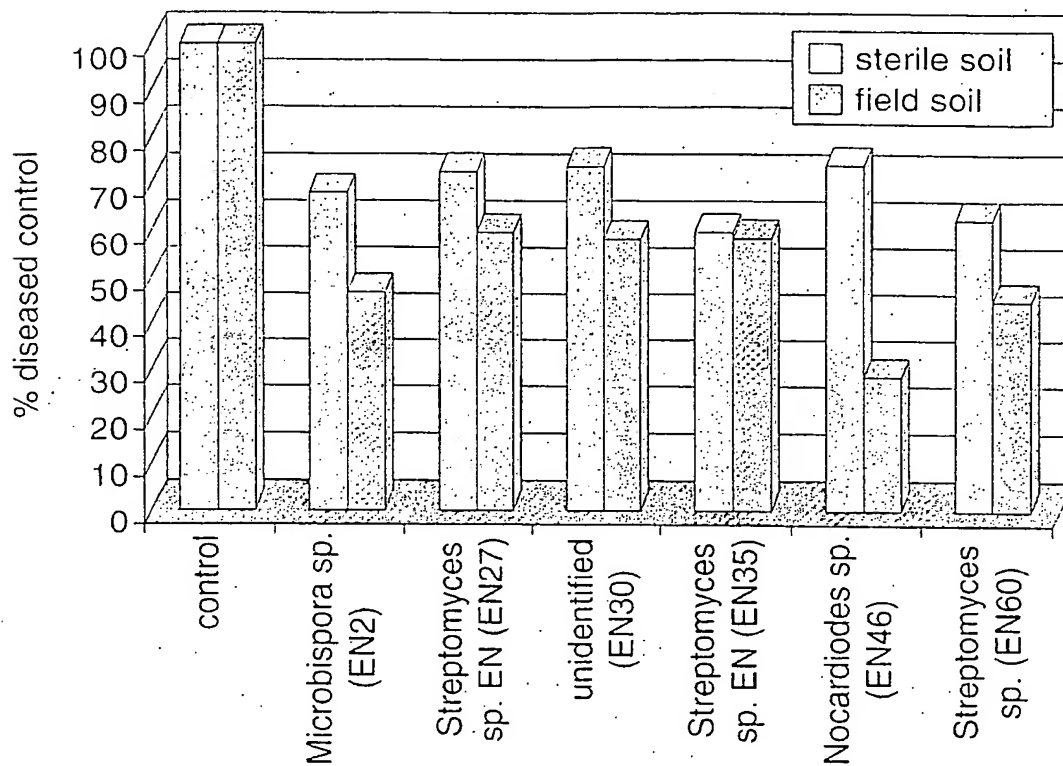


Figure 1

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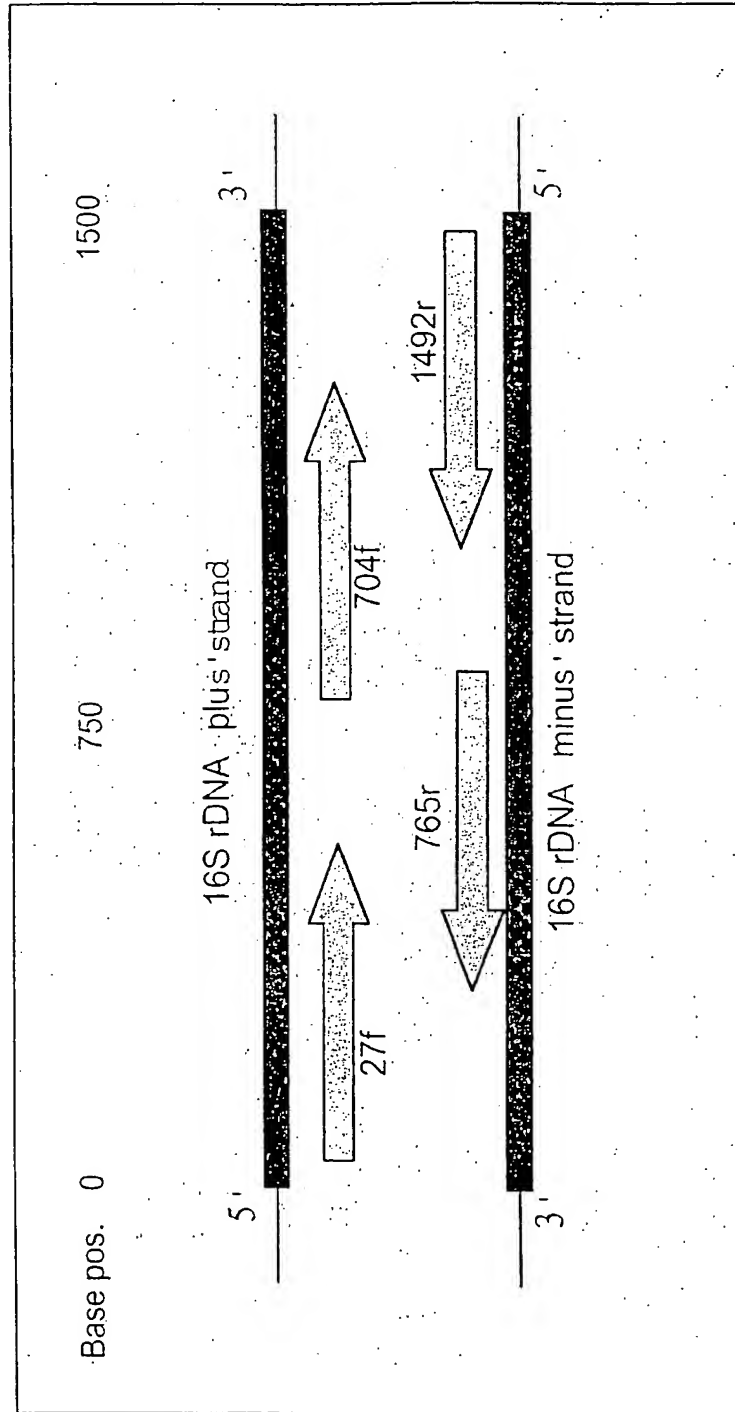


Figure 2

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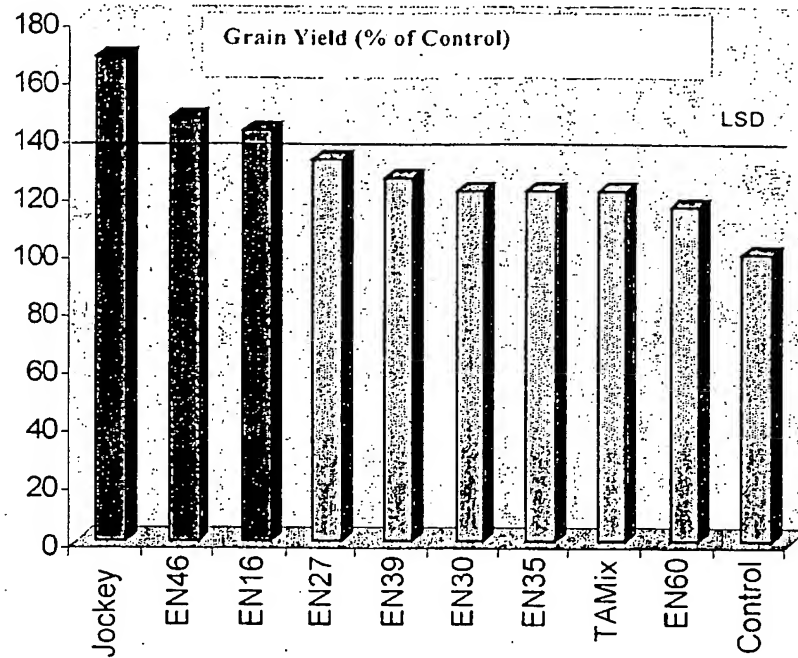


Figure 3

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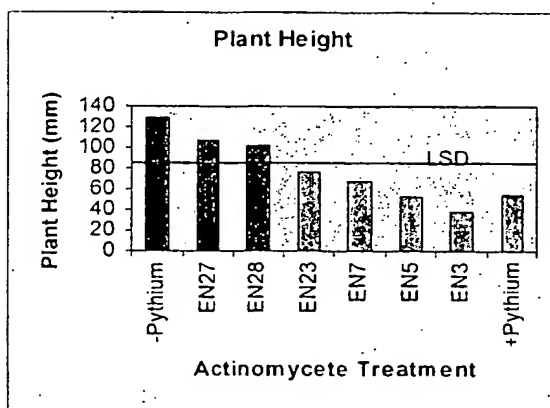


Figure 4

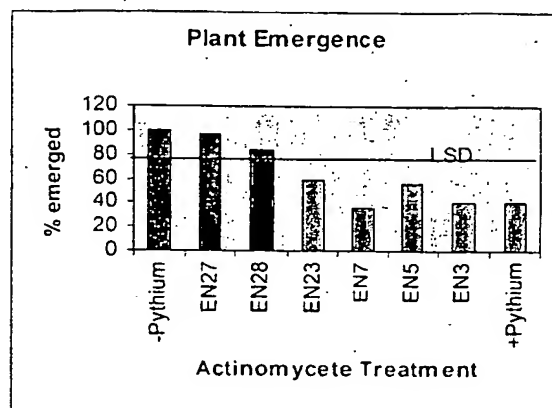


Figure 5

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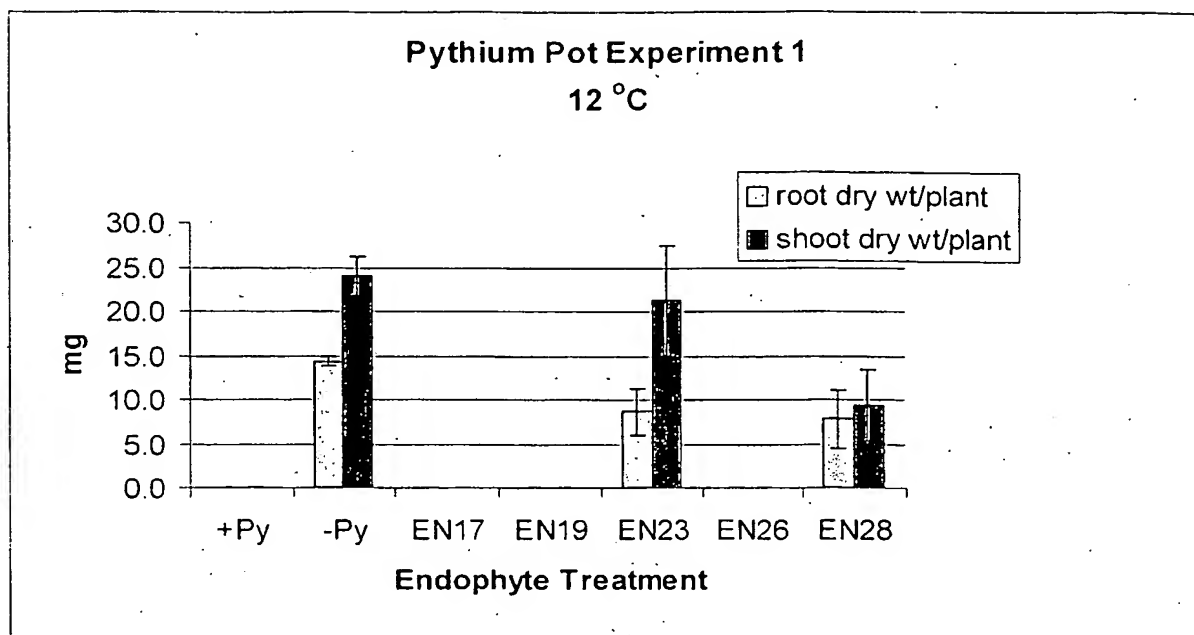


Figure 6

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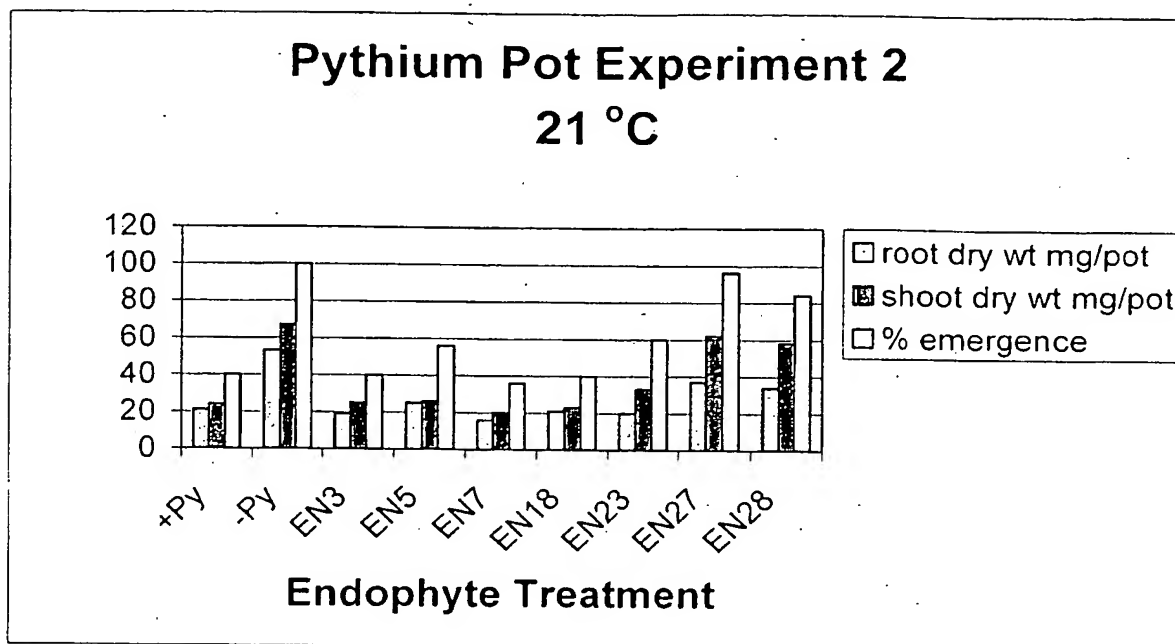
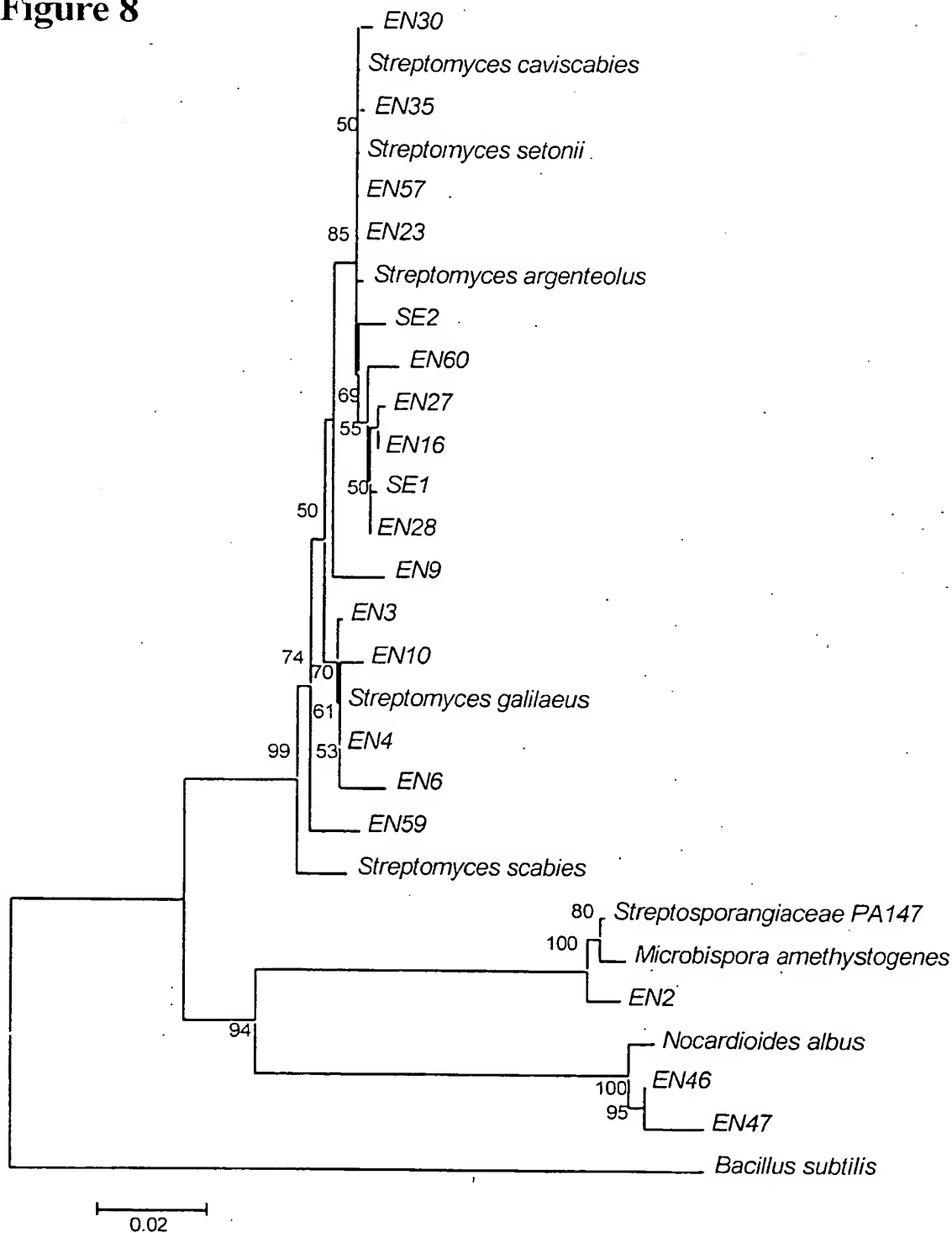


Figure 7

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Figure 8



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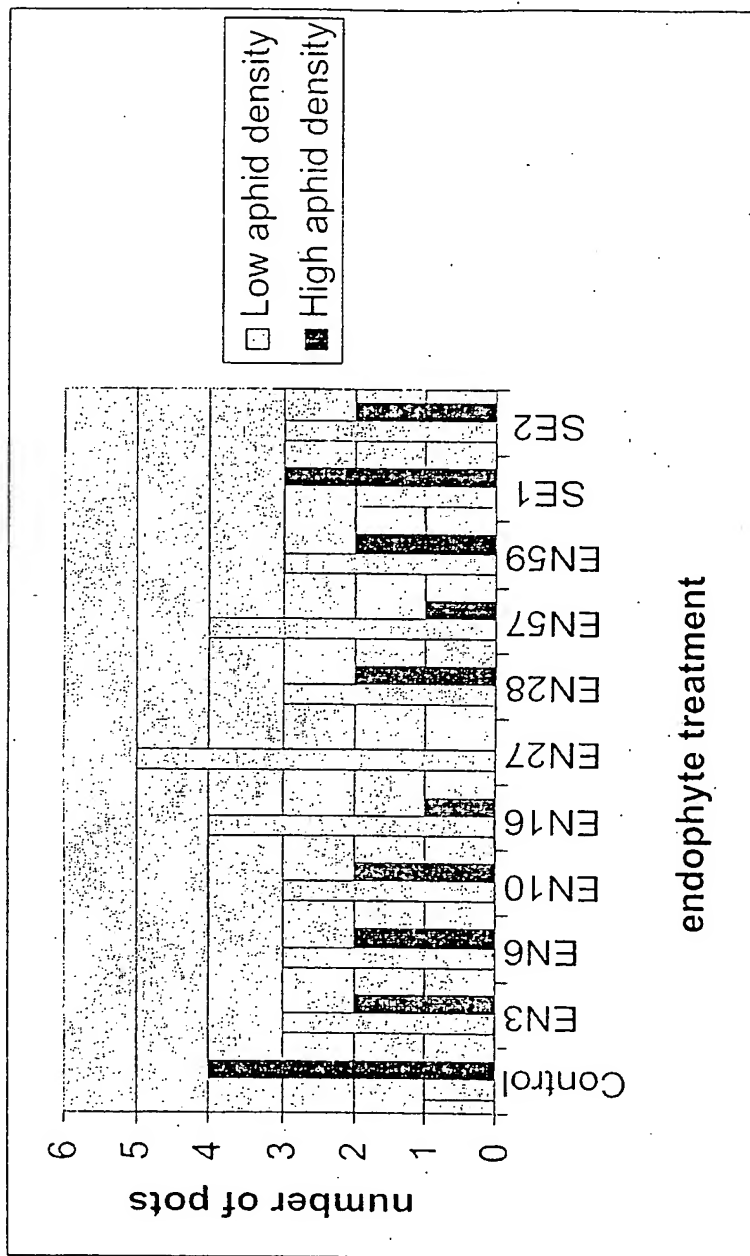


Figure 9

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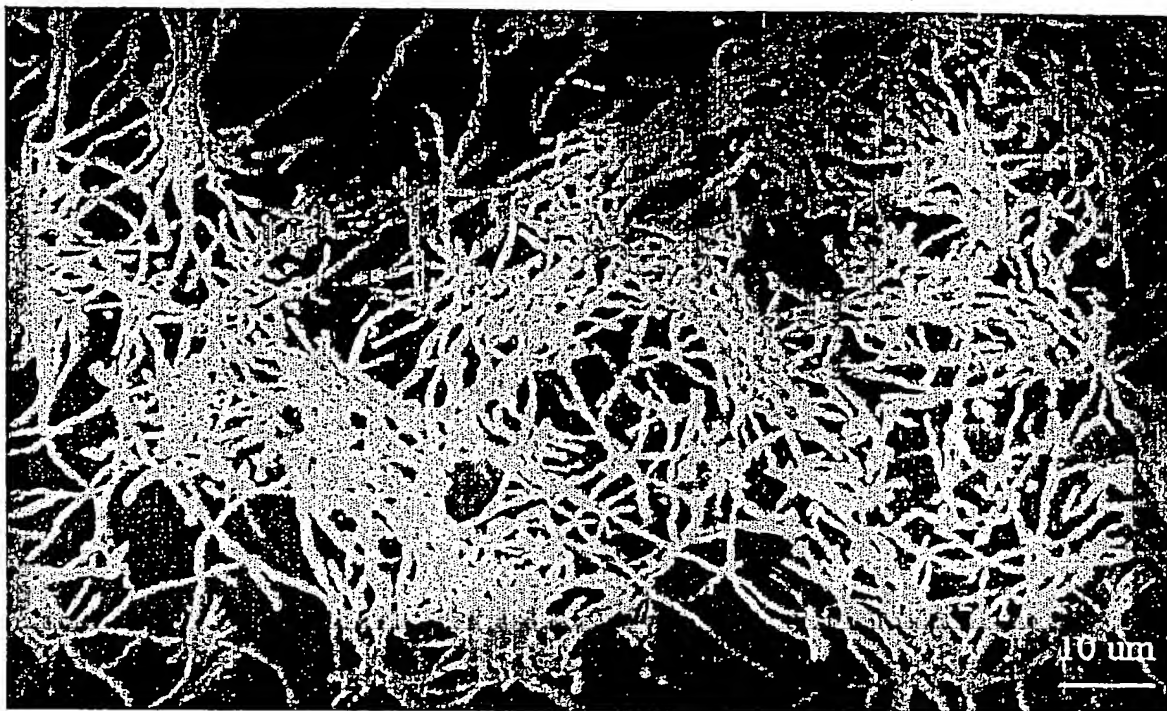
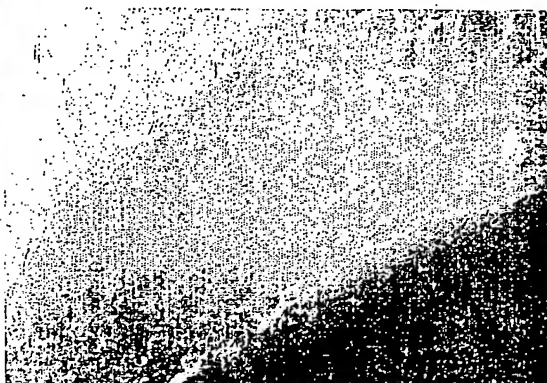


Figure 10

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11.1 Blue excitation/ green emission

11.2 UV excitation/ blue emission



11.3 Image enhanced merge



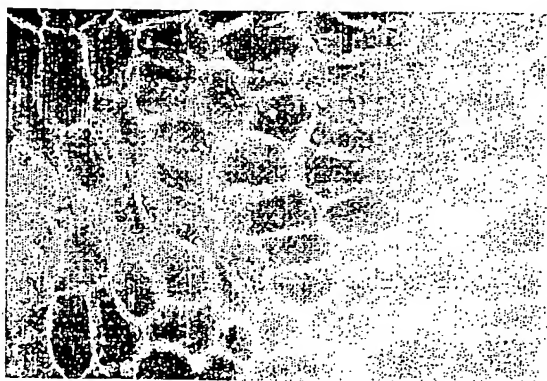
Figure 11

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12.1 Blue excitation/ green emission



12.2 UV excitation/ blue emission



12.3 Image enhanced merge

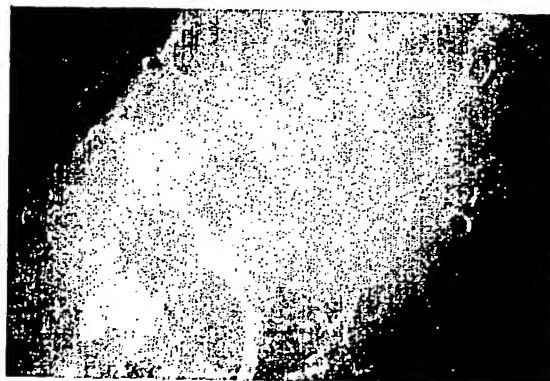


Figure 12

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13.1 Blue excitation/ green emission

13.2 UV excitation/ blue emission



13.3 Image enhanced merge

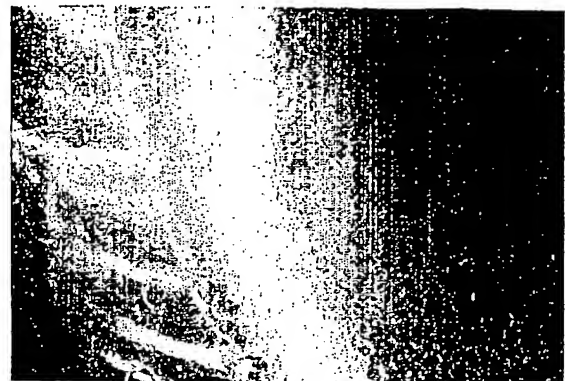
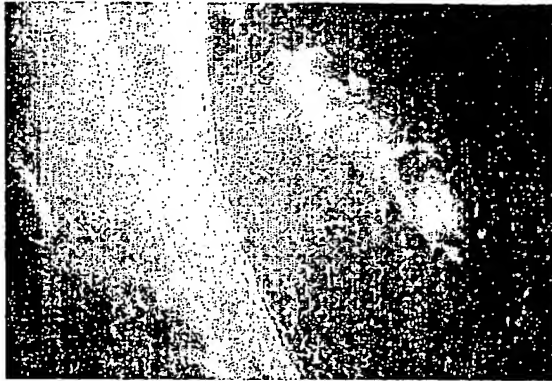


Figure 13

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14.1 Blue excitation/ green emission

14.2 UV excitation/ blue emission



14.3 Image enhanced merge

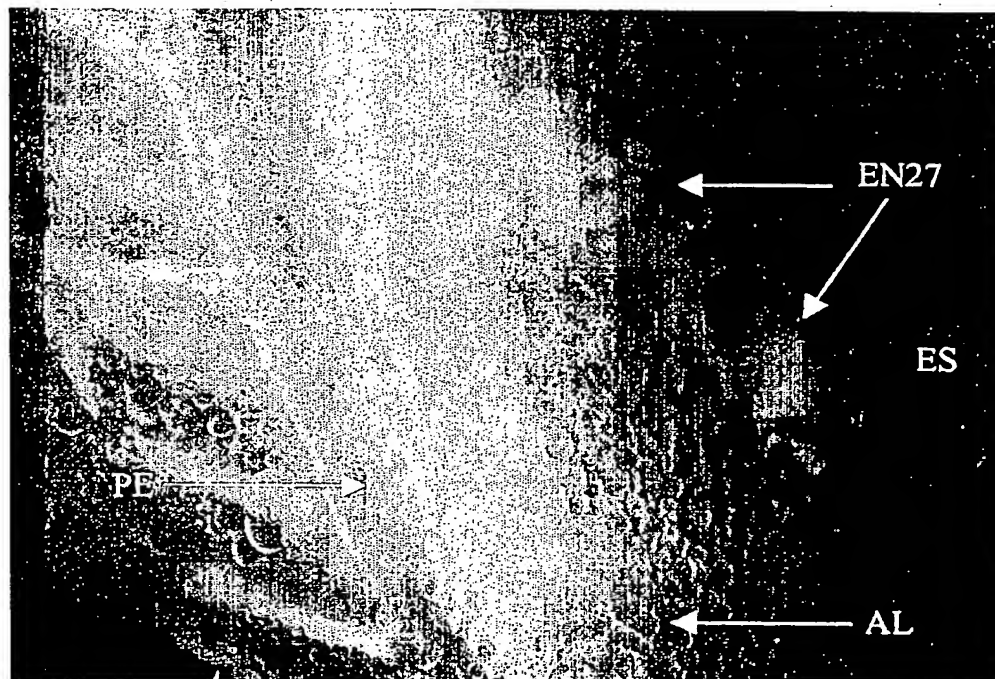


Figure 14

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FIGURE 15

EN2

SEQ ID NO:1

CTTAACACATGCAANTCAAGCGGAAAGGCCCTTCGGGGTACTCAANCGGCNAACGGGTGATTAACACNTGANTAA
CCTGCCCCCTGACTCTGGGATAANCCCTGGGAACTGGGTCTAATACCGGATACAACCATTCTCNCATGGGATGGT
GGTGGAANTTTTTNCGGTTGGGGATGGGCTCGCGGCTATCACCTTGTGGTGGGGTGTAGGCCTACCAAGGCg
ACgAACGGTAgCCCCCTGAgAGGGCgACCGGCCaCaCTGGGACTGAgACaCCGCCgAACTCCTaCgGGAGGCA
gCACTgGGGAATaTTGCCATGGGCGGAAGCCTGACGCAGNACGCCGCTGGGGGATGACGGCCTTNGGGTTGT
AAACCTN'TTTCAGCAGGGACGAAGTTGACGTGTACCTGTAGAAGAAGCGCCGGCTAAATANGTGCCAGCAGCCGC
GGTAATANGTAGGGCGCGAGCGTTNTCCGGAATTATTGGGCGTAAAGAGTTTGTAGGTGGCTTGTTCGTTTGCC
GTGAAAGCCCCGTGGCTTAANTACGGGTTTGGCGGTGGATACGGGCAGGCTAGAGGCTGGTAGGGGCAAGCGGAATT
CCTGGTGTAGCGGTGAAATGCGCAGATATCAGGAGGAACACCGGTGGCGAAGGCGGCTTGTGGGCCAgTTCtGA
CGGTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAGATACCCCTGGTAGTCCACGCTGTAAACGTTGGGCGCTA
GGTGTGGGGTCTTCCACGATCTCTGTGCCGTAGCTAACGCATTAAGCGCCCCGCTGGGGAGTACGGCCGCAAG
GCTAAAACTCAAAGGAATTGACGGGGGGCCCGCACAAAGCGCGGAGCATGTTGCTTAATTGACGCAACGCGAAGA
ACCTTACCAAGGTTTGACATACACCGGAAACACTCANANATGGGTGCCTCCTTTGGACTGGGTACAGGTGGTGC
ATGGCTGTcNNACCCCTCGTGTcGTNAGATGTNGGGTTAAGTCCCgCAACGANCGCAACCCTTGGTTCCATGTTG
CCAGCACNCCCTTTGNGGTGGTGGGGACNCATGGGANAATGCCGGGGTcNACTCNGGAGGAAGGTGGGGATGACC
TCAAGTNATcNTGCCCCCTTATGTTCTTGNNGTG

EN3

SEQ ID NO:2

GCTGGCGGCGTGCTTAACACATGCAAGTGAACGATGAACCACTTCGGTGGGGATTAGTGGCGAACGGGTGAGTA
ACACGTGGGCAATCTGCCCTTCACTCTGGGACAAGCCCTGGAAACGGGGTCTAATACCGGATAACACTNCTGCTC
TCATGGGCAGGGGTTAAAAGCTCCGCGGCTGAAGGATGAGCCCGCGGCCTATCAGCTTGTGGTGAAGTAATGGC
TACCAAGGCGACGACGGGTAGCCGGCCTGAGAGGGCGACCGGCCACACTGGGACTGAGACACGGGCCAGACTCC
TACGGGAGGCAGCAGTGGGGAATATTGCaCaATGGGCGAAAGCCTGATGCAGCGACCGCGCTGAGGGATGACG
GCCTTCGGGTTGTAAACCTCTTTCAGCAGGGAAGAAGCGAAAGTGACGGTACCTGCAGAAGAAGCGCCGGCTAAC
TACGTGCCAGCAGCCGCGGTAATACGTAgGGCGCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGCTtGTAgGC
GGCTTGTACGTCGGGTGTGAAAGCCCCGGGGCTTAACCCcGGGTCTGCATTGATACGGGCTAgCTAgAGTGTGG
TAGGGGAGATCGGAATTcTGGTGTAGCGGTGAAATGCgCAGATATCAGGAGGAACACCGGTGGCGAAGGCGGaT
cTcTGGGCCATTACTGACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAGATACCCCTGGTAGTCCACGCC
GTAAACGGTGGGAAGTAgGTGTTGGCGACATTCCACGTCTGCGGTGCCGAGCTAACGCATTAAGTTCCCCGCCT
GGGGAGTACGGCCGCAAGGCTAAAACCTCAAAGGAATTGACGGGGGGCCCGCACAAAGCAGCGGAGCATGTGGCTTAA
TTCGACGCAACGCGAAGAACCTTACCAAGGCTTGACATACACCGGAAAGCATCAGAGATGGTGGCCCCCTTGTGG
TtCGGTGTACAGGTGGTGCATGGCTGTCTGCTCAGCTCGTGTCTGAGATGTTGGGTAAAGTCCCGCAACGAGCGCA
ACCCTTGTtCTGTGTTGCCAGCATGCCCTTCGGGGTGTAGGGGACTCACAGGAGACCGCCGGGGTCAACTCGGAG
GAAGGTGGGGACGACGTCAAGTCATCATGCCCTTATGTCTTGGGCTGCACACGTGCTACAATGGcCGGTACAAA
GAGCTGCGATACCGTGAGGTGGAGCGAATCTCAAAAAGCCGGTCTCagTTCCGGATTGGGGTcTGCAACTCGAcCC
CATGAAGTCGGAGTTGcTAATAATCgCANATCagCATTGCTGCGGTGAATACGTTcCCGGGCCTTGTAACcACCG
CCcGTACGTcACGAAAGTCGgTAAcACCCgAAgCCGGTGGCCAACCCCTTgTGGGAGGgAGCTGTCTGAAGGTGG
GACTGGCGATTG

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[FIGURE 15 CONTINUED]

EN16

SEQ ID NO:7

GCTTNTTGGTGGGNCNATGGCCTACCAAGGNGAGGACGGNTANCCNGCCTGNGAGGGAGACCGNCCACACTGGGA
ATGNGANACGGCCCAGAATCCTACGGGAGGCAGCANNGGGGAANATTGCACAANGGGCGAAAGCCTGATGCAGNG
ANGCCGCGTGAGGGAAGACGGCCTTTGGGTTGTAAACCTNTTTNAGCAGGGAAGAAGCGAAAGTGACGGTACCTG
CAGAAGAAGCGCCGGCTAANTANGTGCCAGCAGCCGCGTAATANGTAGGGCGCAAGCGTTGTCCGGAATTATTG
GGCGTAAAGAGCTTGTAGGCGGCTTGTGANGTNGGATGTGAAAAGCCCGGGCTTAACCCCGGGTTTGCATTGTAT
ACGGGCTAGCTAGAGTGTGGTAGGGGAGATNGGAATTCCTGGTGTAGCGGTGAAATGCGCAGATATCAGGAGGAA
CACCGGTGGCGAAGGCGGATCTCTGGGCCATTACTGACGCTGAGGAGCGAAAGCCTGGGGAGCGAACAGGATTAG
ATACCTTGGTAGTCCACGCCGTAAACGTTGGGAACCTAGGTGTTGGCGACATTCCACGTCGTCGGTGCCGCAGCTA
ACGCATTAAAGTTCCCCGCTGGGGAGTACGGCCGCAAGGCTAAAACCTCAAAGGAATTGACGGGGGCCCCGCACAAG
CAGCGGAGCATGTGGCTTAATTGACGCAACGCGAAGAACCCTTACCAAGGCTTGACATATACCGGAAAAGCATCAG
AGATGGTGCCCCCTTGTGGTTCGGTATACAGGTGGTGCATGGCTGTCGTCAGCTCGTGTGCTGAGATGTTGGGTT
AAGTCCCGCAACGAGCGCAACCCTTGTCTGTGTTGCCAGCATGCCCTTCGGGGTGATGGGGACTCACAGGAGAC
TGGGGGTCTGNAACCTGACCCCATGAANTCGGAGTTGCTAATAATCCCAAATTTCANCATTGGTGCGGTGAATACT
TCCCCGGCCTGGTACACNACCGCCCGTCAACTCACGAAAGTCGGTNAAACCCGAAACCGGTGGGCCAACCCCTTG
TGGGAAGGAACTGGCCNAAGTGGGACTGGCGATTGGGAC

EN23

SEQ ID NO:10

ACGAACGCTGGCGGCGTGCTTAACACATGCAAGTGAACGATGAAGCCGCTTCGGTGGTGGATTAGTGGCGAACG
GGTGAGTAACACGTGGGCAATCTGCCCTTCACTCTGGGACAAGCCCTGGAAACGGGGTCTAATACCGGATAACAC
TCTGTCCCGCATGGGACGGGGTTGAAAGCTCCGGCGGTGAAGGATGAGCCCGCGGCCTATCAGCTTGTGGTGGG
GTAATGGCCTACCAAGGCGACGACGGGTAGCCGGCCTGAGAGGGCGACCGGCCACACTGGGACTGAGACACGGCC
CAGACTCCTACGGGAGGCAGCAGTGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAGCGACGCCGCGTGAGG
GATGACGGCCTTCGGGTTGTAAACCTCTTTCAGCAGGGGAAGAAGCGAAAGTGACGGTACCTGCAGAAAGAAGCGCC
GGCTAACTACGTGCCAGCAGCCGCGGTAATACGTAGGGCGCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGCT
CGTAGGCGGCTTGTACGTCGGATGTGAAAGCCCGGGGCTTAACCCCGGTCTGCATTGATACGGGCTAGCTAG
AGTGTGGTAGGGGAGATCGGAATTCCTGGTGTAAGCGGTGAAATGCGCAGATATCAGGAGGAACACCGGTGGCGAA
GGCGGATCTCTGGGCCATTACTGACGTCTGAGGAGCGAAAGCGTGGGgAgCGAACAGGATTAGATACCCTGgTAG
TCCACGCCGTAAACGTTGGgAACTAGgTGTTGGCGACATTCCACGTCGTCGGTGCCGCAGCTAACGCATTAAGTT
CCCCGCTTGGGAGTACGGCCGCAAGGCTAAAACCTCAAAGGAATTGACGGGGGCCCGCACAAAGCAGCGGAGCATG
TGGCTTAATTGACGCAACGCGAAGAACCCTTACCAAGGCTTGACATATACCGGAAAGCATCAGAGATGGTGCCCC
CCTTGTGGTTCGGTATACAGGTGGTGCATGGCTGTGCTCAGCTCGTGTGCTGAGATGTTGGGTTAAGTCCCGCAAC
GAGCGCAACCCTTGTCTGTGTTGCCAGCATGCCCTTCGGGGTGATGGGGACTCACAGGAGACTGCCGGGGTCAA
CTCGGAGGAAGGTGGGGACGACGTCAAGTCATCATGCCCTTATGTCTTGGGCTGCACACGTGCTACAATGGCCG
GTACAATGAGCTGCGATGCCGCGAGGCGGAGCGAATCTCAAAAAGCCGGTCTCAGTTCGGATTGGGGTCTGCAAC
TCGACCCCATGAAGTCGGAGTTGCTAGTAATCGCAGATCAGCATTGCTGCGGTGAATACGTTCCCGGGCCTTGTA
CACACCGCCCGTCACGTCACGAAAGTCGGTAACACCCGAAGCCGGTGGCCCAACCCTTGTGGGAGGGAGCTGTCTG
AAGGTGGGACTGGCGATTG

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[FIGURE 15 CONTINUED]

EN27

SEQ ID NO: 12

TTAANACATGCAANTCGAACGATGAACCCNGTTTCGGTGGTGGATTAGTGGCGAACGGTGAGTAANANGTGGGCA
ATTTGCCCTTCATTTTGGACAAGCCCTGGAAACGGGTTTAATACCGGATAACATTTTNTCCCGCATGGGANGGGG
TTGAAAGNTCCGGCGGTGAAGGATGAGCCCGCGGCTATNAGCTTGTGGTGGGGTAATGGCCTACCCAAGGGAG
ACGGGTAGCCGGCCTGAGAGGGCGACCGGCCACACTGGGAATGAGANACGGCCCAGAATCCTACGGGAGGCAGCA
GTGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAGCGANGCCGCGTGAGGGATGACGGCCTTNGGGTTGTAA
ACCTTTTNNAGCAGGGAAGAAGCGAAAGTGACGGTACCTGCAGAAGAAGCGCCGGCTAAATAAGTGCCAGCAGCC
GCGGTAATAAGTAGGGCGCAAGCGTTGTCCGAATTATTGGGCGTAAAGAGCTTGTAGGCGGCTTGTCANGTNGG
ATGTGAAAGCCCCGGGNTTAACCCCGGGTTTGCATTTGATACGGGCTAGNTAGAGTGTGGTAGGGGAGATNGGAA
TTCCTGGTGTAGCGGTGAAATGCGCAGATATCAGGAGGAACACCGGTGGCGAAGGCGGATCTCTGGGCCATTACT
GACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAGATACCCTGGTAGTCCACGCCGTAACCGTTGGGAAC
TAGGTGTTGGCGACATTCCACGTCGTGCGTGCCGACGCTAACGCATTAAGTTCCCGNCTGGGGAGTACGGCCGC
AAGGCTAANACTCAAAGGAATTGACGGGGGCCCCGNACAAGCAGCGGANCATGTGGCTTAATTCGACGCANCGCGA
AGAACCTTACCAAGGCTTGACATATACCGGAAAGCATCAGAGATGGTGCCCCCTTGTGGTGCNTATACANGTGG
TGCATGNTGTGTCACCTCGTGTGTCGTGAGATGTTGGGTTAAGTCCCGCAACGAGCGCNACCCTTGNTCTGTGTT
GNCANCATGCCCTTCGGGGNTGATGGGGACTCACAGGANAAGTGNCCGGGTCAACTCCGGANGAAGGTGGGTGAC
GAAGTCAAGGTATCATGNCCCCTTATGTCTTGGTGCTGCACACGTGC

EN28

SEQ ID NO: 13

TTCGGNGGTGGANTAGNGGCGNACGGGNGACCAACANGNGGGCAATCCCCCTTCANTTTNGGACAACCCCTGGA
AACGGGTTNTAATACCGGATAACANTTTNTCCCGCATGGGANGGGGTTGAAAGCTCCGGCGGTGAAGGATGAGC
CCGCGGCCTATCAGCTTGTGGTGGGGTAATGGCCTACCAAGGCGACGACGGGTAGCCGGCCTGAGAGGGCGACC
GGCCACACTGGGANTGAGANACGGCCCAGAATCCTACGGGAGGCAGCAGTGGGGAATATTGCACAATGGGCGAAA
GCCTGATGCAGCGACCGCGGTGAGGGATGACGGCCTTCGGGTTGTAAACCTTTTTCAGCAGGGAAGAAGCGAAA
GTGACGGTACCTGCAGAAGAAGCGCCGGCTAAATANGTGCCAGCAGCCGCGGTAATANGTAGGGCGCAAGCGTTG
TCCGGAATTATTGGGCGTAAAGAGNTTGTAGGCGGCTTGTGANGTCGGATGTGAAAGCCCCGGGCTTAACCCCGG
GTTTGCATTCGATACGGGCTAGCTAGAGTGTGGTAGGGGAGATCGGAATTCCTGGTGTAGCGGTGAAATGCGCAG
ATATCAGGAGGAACACCGGTGGCGAAGGCGGATcTcTGGGCCATTACTGACGcTGAGGAGCGAAAGCGTGGGgAG
CGAACAGgAATTAGATACCCTGgTAGTCCACGCCGTAACGTTGGgAAcTAGgTGTGGcGACATTCCACGTcGT
CGgTGCCCGCAGCTAACGCATTAAGTTCCCGCCTGGGGAGTACGgCCCGCAAGGCTAAACTCAAAGGAATTGAC
GGGGgCCCGCACAAAGCAGCGGAGCATGTGGCTTAATTCGACGCAACGCGAAGAACCTTACCAAGGCTTGACATAT

ACCGGAAAGCATCAGAGATGGTGCCCCCTTGTGGTGGTATACAGGTGGTGATGGCTGTGTCGTGAGCTCGTGTC
GTGAGATGTTGGGTTAAGTCCCGCAACGAGCGCAACCCTTGgTTCTGTGTTGgCCAGCATGCCCTTCGGGGTGAT
GGGGACTCACAGGAGACTGgCCGGGGTCAACTCGGAGGAAGGTGGGGACGACGTCAAGTCATCATGCCCTTATG
TCTTGGGgCTGCACACGTGCTACAATGGCCGGTACAATGAGCTGCGATGCCGCGAaGGCGGAGCGAATCTCAAAA
AaGCCGGTCTCAGTTCCGATTGGGGTCTGCAACTCGACCCCATGAAGTCGGAGTTGCTAGTAATCGCAGATCAGC
ATTGCTGCGGTGAATACGTTCCCGGGCCTTGACACACCGCCCGTCACGTACGAAAGTCGGTAACACCCGAAGC
CGGTGGTCCAACCCCTTGTGGGAGGGAGCTGTGCAAGGTGGGACTGGCGATTGG

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[FIGURE 15 CONTINUED]

EN46

SEQ ID NO:16

ATGCAAGTCGAGCGGAAAGGCCCTTCGGGGTACTCGAGCGGCGAACGGGTGAGTAACACGTGAGTTAATCTGCCC
CAGGCTCTGGATACCCACCGGAAAACGGTGATTAATACCGAATACGACAACCGATTTGCATGATCTGGTgGTGNA
AAGTTTTTCGGCCTGGGATGTGCTTCGCGGCCTATCAGCTTGTTGGTGAGGTAATGGCTCACCCAAGGCTTCGAC
GGTAGCCGGCCTGAGAGGGTGACCGNCCACACTGGGACTGAGACACGGCCCAGACTCCTACGGGAGGCAGCAGTG
GGGAATATTGGACAATGGGCGGAAGCCTGATCCAGCAACGCCGCGTGAGGGATGACGGCCTTCGGGTTGTAAACC
TCTTTCAGCACAGACGAAGCGCAAGTGACGGTATGTGCAGAAGAAGGACCGGCAACTACGTGCCAGCAGCCGCG
GTAATACGTAGGGTCCGAGCGTTGTCCGGAATTATTGGGCGTAAAGGGCTCGTAGGCGGTCTGTCCGCTCGGGAG
TGAAACCAGGTGCTTAACACCTGGCCTGCTTTCGATACGGGCAGNCTAGAGGTACNCAGGGGAGAATGGAATTC
CTGGTGTAGCGGTGAAATGCGCAGATATCAGGAGGAAACACCGGTGGCGAAGNCGGTTCTCTGGGAGTATCCTGA
CGCTGAGGAGCGAAAGTGTGGGGAGCGAACAGGATTAGATACCCTGGTAGTCCACACCGTAAACGTTGGGCGCTA
GGTGTGGGACACATTCCACGTGTTCCGTGCCGCAGCTAACGCATTAANCGCCCCGCTGGGGAGTACGGCCGCAA
NGCTAAACTCANAGGAATTGACGGGGGGCCCGCACAAAGCGGCGGA_gCATGCGGATTAATTCGATGCAACGCGAAG
AACCTTACCTGGGTTTGACATACACCGGAAAGCCGTACAGATACGGCCCCCTTTTAGTC_gGTGTaCAGGTGGTGCA
TGGCTGTCT_gCa_gCt_gCTGTCTGTGAGATGTtCGGGTTAAGTCCCGCAACGAGCGCAaCCCTC_gTCCTATGTTGC
CaGCAATTTCGGTTGgGGACTCATAGGA_gACTGCC_gGGGTCaACTCGGAGGAAGGTGGGGATGACGTCAAGTCATC
ATGCCCTTATGTCCAGGGCTTCACGCATGCTACAATGGCCGGTACAAAGGGCTCGCATCCCGTGAGGGTGAGCG
AATCCCAAAAAGCCGGTCTCAGTTCGGATTGGGGTCTGCAACTCGACCCCATGAAGTCGGAGTCGCTAGTAATCG
CAGATCAGCAACGCTGCGGTGAATACGTTCCCGGGCCTTGTTACACACCGCCCGTCACGTACGAAAGTCGGCAAC
ACCGAAGCCANTGGCCCAACTCGTAAGAGAGGGAGCTGT

EN60

SEQ ID NO:18

ATGCAAGTNGAACGATGAANCCNTTTGGGGTGGATTAGTGGCGAACGGGTGAGTAANANGTGGGCAATTTGCCCT
TCAATTTGGGGAACAGCCCTGGAAACGGGGTNTAATACCGGATAACANTNTGTCCCGCATGGGACGGGGTTAAAG
CTCCGGCGGTGAAGGATGAGCCCGCGGCCTATNAGCTTGTTGGTGGGTGATGGCCTACCAAGGCGACGACGGGT
AGCCGGCCTGAGAGGGCGACCGGCCACACTGGGACTGAGACACGGCCAGACTCCTACGGGAGGCAGCAGTGGGG
AATATTGCACAATGGGCGAAAGCCTGATGCAGCGACGCCGCGTGAGGGATGACGGCCTTCGGGTTGTAAACCTTT
TTCAGCAGGGAAGAAGCGAAAGTGACGGTACCTGCAGAAGAAGCGCCGGCTAAATANGTGCCAGCAGCCGCGGTA
ATANGTAGGGCGCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGTTTGTAGGCGGCTTGTACGTTNGGATGTGA
AAGCCCGGGGCTTAACCCCGGGTTTGCAATTCGATACGGGCTAGCTAGAGTGTGGTAGGGGAGATCGGAATTCCTG
GTGTAGCGGTGAAATGCGCAGATATCAGGAGGAACACCGGTGGCGAAGGCGGaTcTcTGGGCCATTACTGACGNT
GAGGAGCGAAAGCGTGGGGAGCNAACAGNATTAGATACCCTGGTAGTCCAAGCCGTAAACGTTGGGAACCTANGTG
TTGGCGACATTCCACGTCTCNNTGCCGCANCTAACGCATTAAGTTCCCCGCCTGGGGAGTACGGCCGCAAGGCT
AANACTCAAAGGAATTGANGNNGCCCGCACAAAGCAGCGGAGCATGTGGCTTANTTCNACGCANCGCGAAGAACC
TTACCAAGGTTTGCCATATAcCGGAAAgCaTCa_gAgATgGTGCCCCCTtGTGGTCGGTATACAGgTGGTGCNTG
GCTGTCTGTCa_gCTCTGTCTGTGACAtGTTGGTTAAgTCCCGTCAaCGAGgCGCAACCCTTGTTNTGTGTNGCCAG
CATGCCCTTCGGGGTGTGGGGACTCACAGGAGACTGCCGGGGTCAACTCGGAGGAAGGTGGGGACGACGTCAAG
TCATCATGCCCCTTATGTCTTGGGCTGCACACGTGCTACAATGGCCGGTACAATGAGCTGCGATGCCGCGAGGCG
GAGCGAATCTCAAAAAGCCGGTNTCAGTTCGGATTGGGGTCTGCAACTCGACCCCATGAAGTCGGAGTTGCTAGT
AATCGCAGATCAGCATTGCTGCGGTGAATACGTTCCCGGGCCTTGTTACACACCGCCCGTCACGTACGAAAGTCG
GTAAACACCCGAAGCCGNTGG

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[FIGURE 15 CONTINUED]

PM87

SEQ ID NO:24

GGCCCAGANATCCGNCTTCGCCACCGGTGTTCCCTCCTGAATATCTGCGCATTTACCCGCTACACCAGGAATTCCG
ATCTCCCCTACCACTCTAACTAGCCCGTATCGAATGCAGACCCGGGGTTAAGCCCCGGGCTTTACATCCGAC
GTGACAAGCCGCCTACGAGCTCTTTACGCCCAATAATTCCGGACAACGCTTGCGCCCTACGTATTACCGCGGCTG
CTGGCAGTAATTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTGCTTCTTCCCTGCTGAAAGAGGTTTACA
ACCCGAAGGCCGTCATCCCTCACGCGGCGTCGCTGCATCAGGCTTTGCCCCATTGTGCAATATTCCCCACTGCTG
NCTCCCGTANGAGTCTGGGCCGTGTCTCAGTCCAGTGTGGCCGGTCGNCCTCTCAGGCCGGCTACCGTCGTGCG
CTTGGTAGGCCATTACCCACCAACAAGCTGATANGCCGNGGGCTCATCCTTCANCGTCGGAGCTTTCAANCCCG
TCCATGCGGGACAGAGTGTTATCCGGTATTANACCCCGTNTCAGGGCTTGTCCANAGTGAAGGGCAGATNGCCAC
GTGTTATCACCGTTCGCCACTAATNACANCGAAACGGCTTATCGTNCGACTGCATGTGTTAACACNCGCAGCGTT
CGTCCTGAGCCAG

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FIGURE 16

EN5.

SEQ ID NO: 3

GTAATGGCCCANAAAACCGCCTTCGCCACCGGTGTTCCCTCCTGATATCTGCGCATTTACCGCTACACCAGGAAT
TCCNATCTCCCCCTACCACACTCTAGCTAGCCCGTATCNAATGCAAACTCGGGGTAAAGCCCNAGCTTTACATC
CGACGTGACAAGCCGCTACAANCTCTTTACGCCCAATAATTCCGGANAACGCTCGCACCCCTACNTNTTACCGCG
GCTGCTGGCNCSTNTTTAGCCGGTGCTTCTTCTGCAGGTACCGTCACTTTCGCTTCTTCCCTGCTNAAAAGGTT
TACAACCTTANGGCCGT

EN6

SEQ ID NO: 4

TGAGGGATGACGGCNTTCGGGGTTGTAAACNTTTNTCACCAGGGAAGAAGCGAAAGTGNCGGTACCTGCAGAAGA
AGCGCCGNTAACTACGGGCCAGCATCCGCGGTAAATACGTAGGGCGCAATCGTTGTCCGGAATTANTGGGCGTAA
AGAGNTCGTAGGCGCTTATCACGTCGGGTGTGAAGCCCCGGGCTTAAGCCCCGGGTCTGCATTTCGATACGGGC
TAGCTAGANTNTGNTAGGGGAGATCGGAATTCCTGGTGTAGCGGTGAAATGCGCAGATATCAGCAGGAACACCGG
TGGCGAAGGCGGATCTCTGGGCCATTACTGACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAGATACCC
TGGTAGTCCACGCCGTAAACGGTGGGAAGTGGTGTGGCGACATTCCACGTCGTCGGTGCCGCAGCTAACGCAT
TAAGTTCCCCGCCTGGGGAGTACGGCCGCAAGGCTAAAACTCAAAGGAATTGACGGGGGCCCCGACAAGCAGCGG
AGCATGTGGCTTAATTCGACGCAACGCGAAGAACCTTACCAAGGCTTGACATACACCGGAAGCATCAGAGATGG
TGCCCCCTTGTGGTGGTGTACAGGTGGTGCATGGCTGTGCTCAGCTCGTGTCTGAGATGTTGGGTTAAGTCC
CGCAACGAGCGCAACCCCTTGGTTCTGTGTGGCAGCATGCCCTTCGGGGTGATGGGGACTCACAGGAGAACGCCG
GGGTCAACTCGGAGGAAGGTGGGGACGACGTCAAGTCATCATGCCCTTATGTCTTGGGCTGCACACGTGCTACA
ATGGCAGGTAAATGAGCTGCGATACCGTGAGGTGGAGCGAATCTCAAAAAGCCTGTCTCANTTCGGATTGGGGT
CTGNAANTCGACCCCATGAAAGTCGGAGTTGCTAATTATCCAGATCAACATTGCTGGCGGTGAATACGTTCCCG
GGGCCTTGGTAAACACCGCCCGTCAANGTNAAGAAAGTCGGGTAACACCCGAAANCCGGTGGGCCAANCCCT

EN7

SEQ ID NO: 5

CCGCCTTCGCCACCGGTGTTCCCTCCTGATATCTGCGCATTTACCGCTACACCAGGAAATTCNATCTCCCCCTA
CCACACTCTANCTANCCCGTATCGAATGCAAAACCGGGGTAAANCCCCGGGCTTTACACCCGACNTGACAAGCC
GCCTACAAACTCTTTACGCCCAATAATCCGGACAACGCTTGCGCCCTACNTATTACCGCGGCTGCTGGCACNTA
TTTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTCGCTTCTCCCTGCTGAAAAAGGTTTACAACCCGAAGGC
CGTCATCCCTCACGCGGCGTCTGCTGCATCAGGCTTTCGCCCATTTGTGCAATATTCCCCACTGCTGCCTCCCTAG
GAATCTGGGCGGTGTCTCAATCCAGTGTGGCCGGTCCCTCTCNGGCCGGCTACCGTCNTCCCTTGGTNACCATT
ANCTACCAACAACCTGATAGGNCGCGGGCTCATCTTCACGCGGGAACCTTCAACCACC

EN9

SEQ ID NO: 6

GGCGGCGTGCTTAACACATGCAAGTCGAACGATGAAGCCCTTCGGGGTGGATTAGTGCGGAACGGGTGAGTAACA
CGTGGGCAATCTGCCCTTCACTCTGGGACAAGCCCTGGAAACGGGGTCTAATACCGGATACGATTCTGAGGCGAT
CTCCTGGTaCTGGAAAGCTCCGGCGGTGAAGGATGAGCCCGCGGCcTATCAGCTTGTTGTGGGTAAATGGCCTACC
AAGGCGACGACGGgTAGCCGGCCTGAGAGGGCGACCGGCCACACTGGGACTGAGACACGGCCAGACTCCTACGG
GAGGCAGCAGTGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAGCGACGCCGCTGAGGGATGACGGCCTTC
GGGTTGTAAACCTCTTTCAGCAGGGAAGAAGCGAGAGTGACGGTACCTGCAGAAGAAGCGCCGGCTAACTACGTG
CCAGCAGCCGCGGTAAATACGTAGGGCGCAAGCGTGTCCGGAATTATTGGGCGTAAAGAGCTCGTAGGCGGCTTG
TCACGTCCGGTGTGAAAGCCCGGGCTTAACCCCGGGTCTGCATCCGATACGGGCAGGCTAGAGTGTGGTAGGGG
AGATCGGAATTCTGGTGTAGCGGTGAAATGCGCAGATATCAGGAGGAACACCGGTGGCGAAGGCGGATCTCTGG
GCCATTACTGACGCTGAGGAgCgAAAGCGTGGGGAGCCaACAGGATTAGATACCCTGGTAGTCCACGCCGTAAAC
GTTGGAAGTGGTGTGGCGACATTCCACGTCTCGGTGCCGACGTAACGCATTAAGTTCCCCGCTGGGGAGT
ACGGCCGCAAGGCTAAAACTCAAAGGAATTGACGGGGGCCCGCACAAGCAGCGAGCATGTGGCTTAATTCGACG
CAACGCGAAGAACCTTACCAAGGCTTGACATATACCGGAAAGCGCCAGAGATGGTGCCCCCTTGTGGTCCGGTAT
ACAGGTGGTGCATGGCTGTGCTCAGCTCGTGTGCTGAGATGTTGGGTTAAGTCCCGCAACGAGCGCAACCCCTGT

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[FIGURE 16 CONTINUED]

CCTGTGTTGCCAGCATGCCCTTCGGGGTGATGGGGACTCACAGGAGACCGCCGGGGTCAACTCGGAGGAAGGTGG
GGACGACGTCAAGTCATCATGCCCTTATGTCTTGGGCTGCACACGTGCTACAATGGCCGGTACAAAGAGCTGCG
ATGCCGTGAGGCGGAGCGAATCTCAAAAAGCCGGTCTCAGTTGGGATTGGGGTCTGCAACTCGACCCCATGAAGT
CGGAGTTGCTAGTAATCGCAGATCAGCATTGCTGCGGTGAATACGTTCCCGGGCCTTGACACACCGCCCGTCA
GTCACGAAAGTCGGTAACACCCGAAGCCGGTGGCCCAACCCCTCGGGAGGGAGCTGTCTGAAGGTGGGAC

EN17

SEQ ID NO:8

CCGCCTTCGCCACCGGTGTTCTCTGATATCTGCGCATTTACCGCTACACCAGGAATTCC
NATCTCCCCTACCACACTCTAGCTAGCCCGTATCAAATGCAAACCCGGGGTTAAGCCCCGGGCTTTC
ACATCCNACGTGACAAGCCGCCTACAANCTCTTTACGCCCAATAATTCCGGACAACGCTTGCGCCCT
ACNTATTACCGCGGCTGCTGGCACNTATTTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTCGCT
NCTTCCCTGCTGAAANAGGTTTACAACCCAAAGGCCNTCATCCCTCNCCGGCNCNTTGCNTCNGGC
TTNCCNCCATTGTTCAANNTTCCCACTGCTNCTCCCTCGGAATCTGGGCCGNTGTCTCATTCCCN
TTNTGGCCGGTCCCCCTCNCAGGCCNGCTACCC

EN19

SEQ ID NO:9

CTCAGCGTCNGTAATGGCCCAAAAACCGCCTTCGCCACCGGTGTTCTCTGATATCTGCGCATTTACCGCTAC
ACCAGGAATTCCNATCTCCCCTACCACACTCTAGCTAGCCCGTATCNAATGCAAACCCGGGGTTAANCCCCGGGC
TTTCACATCCNACNTGACAAGCCGCCTACAANCTCTTTACGCCCAATAATTCCGGACAACGCTTGCNCCCTACTT
ATTACCGCGGCTGCTGGCACTTATTTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTCGCTTCTTCCCTGCTN
AAAAAGGTTTACAACCCNAAGGCCGTCATCCCTCACGCGCNCCTGCTGCATCAGGCTTTCNCCCATTTGTGCAATA
TTCCCCACTGCTGCCTCCCGTAGGATTCTGGGCCGNTCTCATTCCANTGTGGCCGGTCCGCTCTCAGGCCGG
CTACCCGTCNCTCNCCTTGGTAGGCCATTACCCACCAACAAGCTNATAGGCCGCGGGCTCATCCTTACCCGCCG
AAGCTTTCAACCCNCTCCATGCGGGANAAATTGTTNTCCGGTATTAAACCCCGTTTCCAGGGNTTGTCCCAAAT
TGAAGGGGGGATTGNCCACTTTTTACTACCCGTTCNCCNCTAATCCACCACC

EN26

SEQ ID NO:11

CCGCCTTCGCCACCGGTGTTCTCTGATATCTGCGCATTTACCGCTACACCAGGAATTCCNATCTCCCCTACC
GAACTCTANCTGCCCCGTATCNACTGCAAACCCGGGGTTAAGCCCCGGGCTTTCACAACCGACNTGACAAGCCGC
CTACAANCTCTTTACNCCCAATAATTCCGGACAACGCTTGCGCCCTACNTATTACCGCGGCTGCTGGCACNTATT
TAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTCGCTTCTTCCCTGCTGAAAAGGTTTACAACCCGAAGCCG
TCNCTCCCTCACGCGCGTGCCTGCATCAGGCTTTCGCCCATTTGTGCAATATTCCCACTGCTGCCTCCCGTAGGA
TTCTGGGCCGTGTCTCANTCCANTNTGGCCGGTCCCTCTCAGGCCGNTACCCGTCGTCCCTTGGTGAACCNC
TACCTCNCAACAANCTGATAGGGCGCGGGCTCANCNTGCACGCCGGANCTT

EN35

SEQ ID NO:14

AACACATGCAAGTCGAACGATGAAGCCGCTTCGGTGGTGGATTAGTGGCGAaCGGGTGAgtAACACGtGGCCAAN
TGTGNCCGTCACTaTGGGACgAAGaCCTTGGAACCGGGGTCTAATACCGGATAACACTCTGTCCCGCATGGGACG
GGGTTGAAAGCTCCGGCGGTGAAGGATGAGCCCGCGGCCTATCAGCTTGTGGTGGGTAATGGCCTACCAAGGC
GACGACGGGTAGCCGGCCTGAGAGGGCGACCGGCCACACTGGGACTGAGACACGGCCAGACTCCTACGGGAGGC
AGCAGTGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAGCGACGCCGCTGAGGGATGACGGCCTTCGGGTT
GTAAACCTCTTTCAGCAGGGAAGAAGCGAAAGTGACGGTACCTGCAGAAGAAGCGCCGGCTAACTACGTGCCAGC
AGCCGCGGTAATACGTAGGGCGCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGCTCGTAGGCGGCTTGTACG
TCGGATGTGAAAGCCCGGGGCTTAACCCCGGGTCTGCATTCGATACGGGCTAGCTAGAGTGTGGTAGGGGAGATC
GGAATTCTTGGTGTAGCGGTGAAATGCGCAGATATTACAGGAGGAACACCGGTGGCGAAGGCGGATCTCTGGGCCA
TTACTGACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTATATACCCTGGTAGTCCACGCCGTAAACGTTG
GGAAGTGGTGTGGCGACATTCCACGTCGTCGGTGCCGACGCTAACGCATTAAGTTCCCGCCTGGGGAGTACG
GCCGCAAGGCTAAAACTCAAAGGAATTGACGGGGGGCCGCACAAGCAGCGGAGCATGTGGCTTAATTTCGACGCAA
CGCGAAGAACCTTACCAAGGCTTGACATATACCGGAAAGCATCAgAGATGGTGCCCCCTTGTGGTGGTATACA

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[FIGURE 16 CONTINUED]

GGTGGTGCATGGCTGTCGTCANCTCGTGTGCTGAGATGTTGGGTTAAGTCCCGCAACGAGCGCAACCCTTGTTCT
GTGTTGCCAGCATGCCCTTCGGGGTGATGGGGACTCACAGGAGACTGCCGgGGTCAaCTCGGAGGAAGGTGGGGA
CGACgTCaAGTCATCATGCCCTTATGTCTTGGGCTGCACACGTGCTACAATGGCCGCTACAATGACCTGCgATG
CCGCGAGGCgGACCGAATCTCAaACAAGCCCGTCTCATTCGGATTGCGGTCTGCaActcCGACCCCATgAAgTCC
GACTtGCTAgTACTCGCACgTCAACaTtGCTGCGCTGaATACgTCCCCGGGCCTTGTACACACCGCCCGTCACGT
CACGAAAGTCGgTAACACCCGAAGCCGGTGGNCCAACCCCTTGTGGGAGGGAGCTGTGCAA

EN39

SEQ ID NO:15

ccgccttcgccaccggtgttcctcctgatatctgcgcatttcaccgctacaccaggaattccnatctcccctacc
acactctagctancccgatcnaatgcaaaccgggggttaacccccgggctttcacaccnactnacaanccgc
ctacaactctttacgccccataattccggacaacgcttgccgctacttattaccgcggtgctggcacttatt
tagccggcgcttcttctgcaggtaccgtcactttcgtctcttccctgctgaaaaagggtttacaacccgaaggng
tcatccctcacgcggcntcgtgcatcaggctttcgccattgtgcaatattccccactgctgcctcccgtagna
ntctgggcccgtntctcantcccagtggtgngcggtcgccctctcaggccggctaccgctcgtcncctnggtnaacc
attanntcaccaacaagctgataggccgcggtcctccttcaccgcccggagcttttaacccctgcccataaaaa
cagangtnttatccggtattanaacccggtttccaggg

EN57

SEQ ID NO:17

GTGCTTAACACATGCAAGTCGAACGATGAAGCCGCTTCGGTGGTGGATTAGTGGCGAACGGGTGAGTAACACGTG
GGCAATCTGCCCTTCACTCTGGGACAAGCCCTGGAAACGGGGTCTAATACCGGATAACACTCTGTCCCGCATGGG
ACGGGGTTGAAAGCTCCGGCGGTGAAGGATGAGCCCGCGGCCTATCAGCTTGTGGTGGGTAAATGGCCTACCAAG
GCGACGACGGGTAGCCGGCTGAGAGGGCGACCGGCCACACTGGGACTGAGACACGGCCCAGACTCCTACGGGAG
GCAGCAGTGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAGCGACGCCGCTGAGGGATGACGGCCTTCGGG
TTGTAAACCTCTTTCAGCAGGGAAGAAGCGAAAGTGACGGTACCTGCAGAAGAAGCGCCGGCTAACTACGTGCCA
GCAGCCGCGGTAAATACGTAGGGCGCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGCTCGTAGGCGGCTTGTCA
CGTCGGATGTGAAAGCCCCGGGGCTTAACCCCGGGTCTGCATTTCGATACGGGCTAGCTAGAGTGTGGTAGGGGAGA
TCGGAATTCTTGGTGTAGCGGTGAAATGCGCAGATATNCAGGAGGAACACCGGTGGCGAAGGCGGATCTCTGGCC
ATTACTGACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAGATACCCTGGTAGTCCACGCCGTAAACGTT
GGGAAGTAGGTGTTGGCGACATTCCACGTCGTGGTGGCGAGCTgAACGCATTAAGTTCCCCGCTGGGGAGTA
CGGCCGCAAGGCTAAAAGCTCAAAGGAATTGACGGGGGCGCACAAAGCAGCGGAGCATGTGGCTTAATTGACGC
AACCGGAAGAACCTTACCAAGGCTTGACATATACCGGAAAGCATCAGAGATGGTGCCCCCTTGTGGTTCGGTATA
CAGGTGGTGCATGGCTGTGCTCAGCTCGTGTGCTGAGATGTTGGGTAAAGTCCCGCAACGAGCGCAACCCTTGTT
CTGTGTTGCCAGCATGCCCTTCGGGGTGATGGGGACTCACAGGAgACTGcCGGGGTCAACTCGGAGGAAGGTGGG
GACGACGTCAAGTCATCATGCCCTTATGTCTTGGGCTGCACACGTGCTACAATGGCCGGTACAATGAGCTGCGA
TGCCgCGAgCGGAgCgAATCTCAAAAAGCCGGTCTCagTTCGGATTGGGGTCTGCAACtCGACCCCATGAAGTC
GGAgtTGCTAgTAATCgCagATCagCATTGCTGcGGTGAATACGTTCCCGGGCCTTGTACACACCGCCGTCAcGT
CACGAAAGTCGGTaACACCCGAAGCCGGTGGCCCAACCgCCTTGTGGGAgGGAAGTTTCCA

GCTGGCGGCGTGCTTAACACATGCAAGTGAACGATGAAGCCGCTTCGGTGGTGGATTAGTGCGAACGGGTGAG
TAACACGTGGGCAATCTGCCCTTCACTCTGGGACAAGCCCTGGAACGGGGTCTAATACCGGATAACACTCTGTCT
CCGCATGGGACGGGGTTGAAAGCTCCGGCGGTGAAGGATGAGCCCGCGGCCTATCAGCTTGTGTGGTGGGTAATG
GCCTACCAAGGCGACGACGGGTAGCCGGCCTGAGAGGGCGACCGCCACACTGGGACTGAGACACGGCCAGACT
CCTACGGGAGGCAGCAGTGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAGCGACGCCGCGTGAGGGATGAC
GGCCTTCGGGTTGTAAACCTCTTTCAGCAGGGAAGAAGCGAAAGTGACGGTACCTGCAGAAGAAGCGCCGGCTAA
CTACGTGCCAGCAGCCGCGETAATACGTAGGGCGCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGCTCGTAgg
CGGCTTGTACGTGCGATGTGAAAGCCCGGGCTTAACCCCGGGTCTGCATTTCGATACGGGCTAgCTAgAGTGTG
GTAGGGGAGATCGGAATTCCTGGTGTAgCGGTGAAATGCGCaGATATCAGGAGGAACAacCGGTGGggAAGgcGGA
TCTCTGGGcCatTACTGACGCTGAGGAGCGAAAAGCGTGGGGAGcGAACAGGATTAGATACCCTGGTAGTCCAAGC
CGTAAACGTTGGGAACTANGTGTGGCGACATTCCACGTCTCGGTGCCGAGCTAACGCATTAAGTTCCCCGTC
CTGGGAGCATACGGCCGCNAGGCTAAAACCTCAAAGGAATTGACGGGGGCCCGCACAAAGcAGcGGAGCATGTGGCTT
ANTTCAGACGCNACCGCAAGAACCTTNCCAAGGCTGACATATACCGgAAAGCATCAcAGATGGTGCCCCCCTTGTG
GTCGGTATACAGGgTGGTGCATGGCTGTtCgtCaGCTCGTGTCgtGAGATGTTGGGTTAaGTCCCGCAAAGAGCG
CAACCgTGTTCTGTGTTGCCAGCATGCCCTTCGGGGTGATGGGGACTCAcAcGAgACTGTcNNGGTCAACTCgga
GGAAGTGGGgGACGACgTCAAGTtCATCATGCCCTTATGCTTGGGCTGCACACNGCTACAATGGCCGGTACA
ATGAGNNGGGATGCCGCGAGGCGGAGCGAATCTCAAAAAcCTCGGTCTCAGTTCGGATTGGGGCTCTGCAACTGACC
CCATGAAGTCGGAGTTGCTAGTAATCGCAGATCAGCATTGCTGCGGTGAATACGTNCCCGGGCCTNGTACACACC
ACCCGTCACGTACGAAAGTCGGTAACACCCTAAGCCGGTGNCCCAACCCCTTNTGGGAGG

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FIGURE 18

PM36

SEQ ID NO:21

CCAGANATCCGCCTTCGCCACCGGTGTTCCCTCCTGATATCTGCGCATTTACCGCTACACCAGGAATTCCGATCT
CCCCTACCACACTCTAGCTAGCCCGTATCGAATGCAGACCCGGGGTTAAGCCCCGGGCTTTCACATCCGACGTGA
CAAGCCGCCTACGAGCTCTTTACGCCCAATAATTCCGGACAACGCTTGCGCCCTACGTATTACCGCGGCTGCTGG
CACGTAGTTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTCGCTTCTTCCCTGCTGAAAGAGGTTTACAACCC
GAAGGNCGTATCCCTCACGCGGCGTGCCTGCATCAGGCTTTCGCCCATTGTGCAATATTCCCCTGCTGCCTC
CCGTAGGAGTCTGGGNCGTGTTCAATNCCAGTGGTGGGCGGTCGCCCTCTCAGGNCGGCTACCGTCTGCGCTT
GGTAGGCATTACCACAACAAGCTGATAGGCGGGGTCATCCTTCAACGCCGGAGCTTCAAACCCGTCCATGCGGG
ACAAGTGTATCCGGTATTAAACCC

PM40

SEQ ID NO:22

TCAGTNATGGCCCAAGANGATCCGNCTTCGCCACCGGTGTTCCCTCCTGATATCTGCGCATTTACCGCTACACCA
GGAATTCCGATCTCCCCTACCACACTCTAACTAGCCCGTATCGAATGCAGACCCGGGGTTAAGCCCCGGGCTTTC
ACATCCGACGTGACAAGCCGCCTACGAGCTCTTNACGCCCAATAATTCCGGACAACGCTTGCGCCCTACGTATTA
CCGCGGCTGCTGGCAGTAGTTAGCCGGCGCTTCTTCTGCAGGTACCGTNACTTTCGCTTCTTCCCTGCTGAAAG
AGGTTTACAACCCGAAGGCCGTCNTCCCTCACGCGGCGTGCCTGCATCAGGCTTTCGCCCCATNGTGCANTATTCC
CCACTGNTGNCTCCCGTANGAGTCTGGGCGGTGTCTCAGTCCCAGTGTGGCCGGTCGNCCTCTCAGGCCGGCTAC
CGTCTGTCGCTTGGTAGGNCATTACCCACCAACAAGCTGATANGTCGNGGGCTCATCCTTACCGNCGGAGNTTT
AACCCCGTNCATGCGGGACAGAGTGTTATCCGGTATTANACCCGTATNCAGGGCTTGTCCCATAGTGAAGGGNAG
ATNGCCACGTGTTATACCGTTCGNCCTAATNATCANCGAANC GGCTTCATCGTTGACTTGCATGTGTTA

PM41

SEQ ID NO:23

CTCAGCGTCAGTCATGGCCAAGAGATCCGCCTTCGCCACCGGTGTTCCCTCCTGTATATCTGCGCATTTACCGCT
ACACCAGGAATTCCGATCTCCCCTACCACACTCTAGCTAGCCCGTATCGAATGCAGACCCGGGGTTAAGCCCCGG
GCTTTCACATCCGACGTGACAAGCCGCCTACGAGCTCTTTACGCCCAATAATTCCGGACAACGCTTGCGCCCTAC
GTATTACCGCGGCTGCTGGCAGTAGTTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTCGCTTCTTCCCTGC
TGAAAGAGGTTTACAACCCGAAGGCCGTCATCCCTCACGCGGCGTGCCTGCATCAGGCTTTCGCCCATTGTGCAA
TATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGGCGGTGTCTCAGTCCCAGTGTGGCCGGTCGCCCTCTCAGGCC
GGCTACCCGTCGTCGCCTTGGTAGGCCATTACCCACCAACAAGCTGATAGGCCGCGGGCTCATCCTTCANCGNCG
GAGCTTTAACCCGTCCATGCGGGACAGAGTGTTATCCGGTATTAAACCCGTTTCAGGGCTTGTCCCANAGTGAAG
GGCAGATTGCCACGTGTTATCANCCGTTTCGNCCTAATCACANCGAANC GGGTTCATCGTTGACTTGCATGTGT
TAA

PM171

SEQ ID NO:25

CCCTCAGGGTCAGTAATGGGCCCAGAGATCCGCCTTCGCCACCGGTGTTCCCTCCTGAATATCTGCGCATTTACCC
GCTACACCAGGAATTCCGATCTCCCCTACCACACTCTAGCTAGCCCGTATCGAATGCAGACCCGGGGTTAAGCCC
CGGGCTTTCACATCCGACGTGACAAGCCGCCTACGAGCTCTTTACGCCCAATAATTCCGGACAACGCTTGCGCCC
TACGTATTACCGCGGCTGCTGGCAGTAGTTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTCGCTTCTTCCC
TGCTGAAAGAGGTTTACAACCCGAAGGCCGTCATCCCTCACGCGGCGTGCCTGCATCAGGCTTTCGCCCATTGTG
CAATATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGGCGGTGTCTCAGTCCCAGTGTGGCCGGTCGCCCTCTCAG
GCCGGCTANCCGTGCTCGCTTGGGTAGGCATTANCCCANCAACAAGCTGATAGGNCGCGGGCTCATNCTTCAAC
GCCGGAGCTTTCAANCCCGTCCATGCGGGACAGAGTGTTATNCGGTATTAAACCCGTTTCAGGGCTTGTTCAGA
GTGAAGGGCAGATTGCCACGTGTTATCAACCGTTCGGCACTAATCACAACGAAGCGGNTTATCGTTGACTTGCA
TGTGTTAACAAGCCGCCAGCGTTTCGTC

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[FIGURE 18 CONTINUED]

PM185-

SEQ ID NO:26

TCAGTAATGGCCCAGAGATCCGCCTTCGCCACCGGTGTTCTCCTGGATATCTGCGCATTTCACCGCTACACCAG
GAATTCCGATCTCCCCTACCACACTCTAGCTAGCCCGTATCGAATGCAGACCCGGGGTTAAGCCCCGGGCTTTCA
CATCCGACGTGACAAGCCGCCTACGAGCTCTTTACGCCCAATAATTCCGGACAACGCTTGCGCCCTACGTATTAC
CGCGGCTGCTGGCACGTAGTTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTTCGCTTCTTCCCTGCTGAAAGA
GGTTTACAACCCGAAGGCCGTCATCCCTCACGCGGCGTCTGCTGCATCAGGCTTTTCGCCCATTGTGCAATATTCCC
CACTGCTGCCTCCCGTAGGAGTCTGGGCGGTGTCTCAGTCCCAGTGTGGCCGGTTCGCCCCTCTCAGGCGGCTACC
CGTCGTCGCCTTGCTAGGCCATTACCCACCAACAAGCTGATAGGCCGCGGGCTCATCCTTCACCGNCGGAGCTT
TAACCCCGTCCCATGCGGGACAGAGTGTATCCGGTATTAGAACCCGTTTCAGGGCTTGTCAGAGTGAAGGG
CAGATTGCCACGTGTTACTCANCCGTTTCGNCACTAATCANCAACGAAGCGGCTTCATCGTTTCGACTTGCATGTGT
TAAGCACGCCGNCAGCGTTCTCTCCTGAGCCAGGATC

PM208

SEQ ID NO:27

TCAGTATCNGCCCAGAGATCCGCCTTCGCCACCGGTGTTCTCCTGATATCTGCGCATTTCACCGCTACACCAG
GAATTCCGATCTCCCCTACCGAACTCTAGCCTGCCCGTATCGACTGCAGACCCGGGGTTAAGCCCCGGGCTTTCA
CAACCGACGTGACAAGCCGCCTACGAGCTCTTTACGCCCAATAATTCCGGACAACGCTTGCGCCCTACGTATTAC
CGCGGCTGCTGGCACGTAGTTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTTCGCTTCTTCCCTGCTGAAAGA
GGTTTACAACCCGAAGGCCGTCATCCCTCACGCGGCGTCTGCTGCATCAGGCTTTTCGCCCATTGTGCAATATTCCC
CACTGGTGNCTCCCGTANGAGTCTGGGGCGGTGTCTCANTCCAGTGTGGGCGGTTCGCCCCTCTCAGGGCGGCTACCGT
CGTCGCTTGCTGAGNCACTACTCACAAACAAGCTGATAGGCCGCGGGCTCATCTGGAACGGCGGAGCTTTACAC

PM228

SEQ ID NO:28

TCAGTAATGGCCCAGANATCCGNCTTCGCCACCGGTGTTCTCCTGATATCTGCGCATTTCACCGCTACACCAGG
AATTCGGATCTCCCCTACCACACTCTAACTAGCCCGTATCGAATGCAGACCCGGGGTTAAGCCCCGGGCTTTCA
ATCCGACGTGACAAGCCGCCTACGAGCTCTTTACGCCCAATAATTCCGGACAACGCTTGCGCCCTACGTATTACC
GCGGCTGCTGGCACGTAGTTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTTCGCTTCTTCCCTGCTGAAAGAG
GTTTACAACCCGAAGGCCGTCATCCCTCACGCGGCGTCTGCTGCATCAGGCTTTTCGCCCATTGTGCAATATTCCCC
ACTGCTGCCTCCCGTANGAGTCTGGGCGGTGTCTCAGTCCCAGTGTGGCCGGTTCGCCCCTCTCAGGCCGGCTACCG
TCGTGCGCTTGCTAGGCCATTACCCACCAACAAGCTGATANGNCGNGGGCTCATCCTTCACCGNCGGAGCTTTCA
ANCCCGTCCCATGCGGGACAGAGTGTATCCGGTATTAAACCCGTTNTCCAGGGCTTGTCATAGTGAAGGGCAGA
TTGCCAAGTGTATCANCCGTTTCGNCACTAATCATCANCAAGCGGCTTCATCGTTTCGACTGCATGTGT

PM252

SEQ ID NO:29

TCCTCAGNATCAGTAATGGCCCAGAGATCCGCCTTCGCCACCGGTGTTCTCCTGATATCTGCGCATTTCACCGC
TACACCAGGAATTCCGATCTCCCCTACCACACTCTANCTAGCCCGTATCGAATGCAGACCCGGGGTTAAGCCCCG
GGCTTTCACATCCGANGTGACAAGCCGCCTACGAGCTCTTTACGCCCAATAATTCCGGACAANGCTTGCGCCCTA
CGTATTACCGCGGNTGCTGGCACGTAGTTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTTCGCTTCTTCCCTG
CTGAAAGAGGTTTACAACCCGAAGGCCGTCATCCCTCACNCGGCGTCTGCTGCATCAGGCTTTTCGCCCATTGTGCA
ATATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGGCGGTGTCTCAATCCCANTGTGGCCGGTTCGCCCCTCTCANGC
CGGCTACCGTTCGCTTGCTAGGCCATTACCCACCAACAAGCTGGATAGGNCGGGGGCTCATCTTCACCGCC
GGAAGCTTTAANCCCGTCCATGCGGGANANAGTGNATCCCNGTATTAAACCCNGTTTCAGGGCTTGTCANAGTG
AAGGGNGATTGCCNAGTGTTTATCNCCCGTTTCGCCANTAATCNACAACGAAGCGGNTTCNTCGNTTCGACTTG
C

25/30

[FIGURE 18 CONTINUED]

PM342

SEQ ID NO:30

TAATGGCCCAGAAATCCGCCTTCGCCACCGGTGTTCCCTCCTGAATATCTGCGCATTTCACCGCTACACCAGGAA
TTCCGATCTCCCCTACCACACTCTAGCTAGCCCGTATCGAATGCAGACCCGGGGTTAAGCCCCGGGCTTTCACAT
CCGACGTGACAAGCCGCCTACGAGCTCTTTACGCCCAATAATTCCGGACAACGCTTGCGCCCTACGTATTACCGC
GGCTGCTGGCACGTAGTTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTCGCTTCTTCCCTGCTGAAAGAGGT
TTACAACCCGAAGGCCGTCATCCCTCACGCGGCGTCTGCTGCATCAGGCTTTCGCCCATTGTGCAATATTCCCCAC
TGCTGCCTCCCGTAGGAGTCTGGGCCGTGTCTCAGTCCCAGTGTGGCGGTGCGCCCTCTCAGGCCGGNTANCCGTC
GTCGCCTTGGTANGCCATTANCCCCACCAACAAGCTGATANGCCGNGGGCTCATCCTTCANCGCCGGAGCTTTTAA
CCCCGTCCCATGCGGGACAGAGTGTTATCCGGTATTAGATCCCGTNTCCAGGGCTTGTCNATAGTGAAGGGCANA
TTGCCACGTGTTACTCANCCGTTCGC

26/30

FIGURE 19

EN4

GGCGGCGTGCTTAACACATGCAAGTCGAACGATGAACCACTTCGGTGGGGATTAGTGGCGAACGGGTGAGTAACA
CGTGGGCAATCTGCCCTTCACTCTGGGACAAGCCCTGGAAACGGGGTCTtAATACCGGATAACACTCCCACCTCTC
CTGAGTGGGGGTAAAAAGCTCCGGCGGTGAAGGATGAGCCCCGGGCCATCAGCTTGTTGGTGAGGTAATGGCTC
ACCAAGGCGACGACGGGTAGCCGGCCTGAGAGGGCGACCGGCCACACTGGGACTGAGACACGGCCCAGACTCCTA
CGGGAGGCAGCAGTGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAGCGACGCCCGGTGAGGGATGACGGCC
TTCGGGTTGTAAACCTCTTTTTCAGCAGGGAAGAAGCGAAAGTGACGGTACCTGCAGAAGAAGCGCCGGCTAACTAC
GTGCCAGCAGCCGCGGTAAATACGTAGGGCGCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGCTCGTAGGGCGGC
TTGTACAGTCCGGGTGTGAAAGCCCGGGCTTAACCCCGGGTCTGCATTTCGATACGGGCTAGCTAGAGTGTGGTAG
GGGAGATCGGAATTCCTGGTGTAGCGGTGAAATGCGCAGATATTACAGGAGGAACACCGGTGGCGAAGGCGGATCT
CTGGGCCATTtACTGACCGTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAGATACCCTGGTAGTCCACGCCGT
AAACGGTGGGAAGTGGTtGTTGGCGACATTCCACGTCTGTCGGTGCCGCAGCTAACGCATTAAGTTCGCCCGCTG
GGGAGTACGGCCGCAAGGCTAAAACTCAAAGGAATTGACGGGGGCCCCGCACAAGCAGCGGAGCATGTGGCTTAAT
TCGACGCAACGCGAAGAACCTTACCAAGGCTTGACATACCCGGAAGCATCAGAGATGGTGCCCCCTTGTGGT
CGGTGTACAGGTGGTGCATGGCTGTCTCAGCTCGTGTCTGAGATGTTGGTTAAGTCCCGCAACGAGCGCAAC
CCTTGTTCTGTGTTGCAGCATGCCCTTCGGGGTGTGAGGACTCACAGGAGACCGCGGGGTCAACTCGGAGGAA
GGTGGGGACgACGTCAAGTCATCATGCCCTTATGTCTTGGGCTGCACACGTGCTACAATGGCAGGTACAATGAG
CTGCGATACCGTGAGGTGGAGCGAATCTCAAAAAGCCTGTCTCAgTTCGGATTGGGGTCTGCAACTCGACCCcaT
GAAGTCGGAGTTGCTAGTAATCGCAGATCAGCATTGCTGCGGTGAATACGTTCCCGGGCCTTGTACACACCGCCC
GTCACGTACGAAAGTCGGTAACACCCGAAGCCGGTGGCCCCAACCCCTTGTGGGAGGGAGCTGTCAAGGTGGGA
CTGGCGATTGGG

EN10

GAGTTTGATCANNGGCTCAGACGAACGCTGGCGGCGTGTTAACACAANCCAAGTCGAANGNTGAACCACTTCGTTG
GGATTAGTGCGAACGGTGNTAACACGNTGGCAATGTGCCCTTCACTNTGGGACAAGNCCTGGAAACGGGGTTCTA
ATACCGGATACCACTACCCGCAGGCATCTGTGGTGTGTTGAAAGCTCCGCCGTGAAGGATGAGCCCCGGGCCAT
CAGCTTGTTGGTGAGGTAATGGCTCACCCAAGGCGACGACGGATAGCCGGCCTGAGAGGGCGACCGGCCACACTG
GGACTGAGACACGGCCAGACTCCTACGGAGGCAGCAGTGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAG
CGACGCCCGGTGAGGGATGACGGCCTTCGGGTTGTAAACCTCTTTTCAGCAGGGAAGAAGCGAAAGTGACGGTACC
TGCAGAAGAAGCGCCGGCTAACTACGTGCCAGCAGCCGCGGTAAATACGTAGGGCGCAAGCGTTGTCCGAATTAT
TGGGCGTAAAGAGCTCGTAGGGCGGCTTGTACGTGCGGTGTGAAAGCCCCGGGGCTTAACCCCGGGTCTGCATTCC
ATACGGGCTAGCTAGAGTGTGGTAGGGGAGATCGGAATTCCTGGTGTACCCGGTGAAATGCGCAGATATCAGGAG
GAACACCGGTGGCGAAGGCGGATCTCTGGGCCATTACTGACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGAT
TAGATACCCTGGTAGTCCACGCCGTAAACGGTGGGAAGTGGTGTGGCGACATTCCACGTCGTGGTGGCCGAG
CTAACGCATTAAGTTCCCCGCTGGGGAGTACGGCCGCAAGGCTAAAACTCAAAGGAATTGACGGGGGCCCCGCAC
AAGCAGCGGAGCATGTGGCTTAATTCGACGCAACGCGAAGAACCTTACCAAGGCTTGACATACGCCGGAAGCAT
CAGAGATGGTGCCCCCTTGTGGTGGTGTACAGGTGGTGCATGGCTGTCTGAGCTCGTGTCTGAGATGTTGG
GTTAAGTCCCGCAACGAGCGCAAcCCTTGTCTGTGTTGcCAGCATGCCCTTCGGGGTGTGAGGACTCACAGGA
GACCGCCGGGGTCAACTCGGAGGAAGGTGGNGACGACGTcAgATCATCATGCCCTTATGTCTTGGGGCTGCACA
CGTGCTACNATGgCaGGTACAATGAGCTGCGATACCGTGAGGTGGAGCgCATCTnnnnnAGCctGTCTCAGTTC
GgATTGGGGTCTGcAACTCGACCCCaTGAAGTCGgAGTTGCTAgATAATCgCAgATCAGCATTGctGCGgTGAAT
ACGttCCCCGGCCTTGTACACACCGCCCTCAGTACGAAAGTCGGTAACACCCGAAGCCGGTGGCCCCAACCC
TTGTGGGAGGGAGCTGTCAANGTGG

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[FIGURE 19 CONTINUED]

EN22

TCCTCAGCGTCAGTAATGGCCCAAAAACCGCCTTCGCCACCGGTGTTCTCCTGATATCTGCGCATTTACCCGCT
ACACCAGGAATTCCNATCTCCCCTACCACACTCTAGCTAGCCCGTATCNAATGCAAACCCGGGTAAANCCCCGG
GCTTTCACATCCNACNTGACAAGCCGCCTACAANCTCTTTACGCCCAATAATTCCGGACAACNCTTGCGCCCTAC
TTATTACCGCGGCTGCTGGCACTTATTTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTTCGCTTCTTCCCTGC
TGAAAAAGGTTTACAACCCNAAGGCCGTCATCCCTCACGCGGCNTCNCCTGCATCAGGCTTTCNCCCATTGTGCAA
TATTCCTCCACTGCTGCCTCCCGTAGGATTCTGGGCCGTNTCTCANTCCANTGTGGCCGGTCGCCCTCTCAGGCC
GGCTACCCGTCGTCNCCTTGGTAGGCCATTACCCCNCCAACAANCTGATAGGCCGCGGGCTCNCCTTCACCGCC
GGAGCTTTCAACCCCGTCCCATGCGGGANAAANTGTTNTCCGGTATTAAAACCCGTTTCCAGGGNTTGTCCAAAA
TTGAAGGGNANATTGCCCACTTTTTNNTCACCCTTCCCCACTAATCCACCACCGAA

EN30

TGGNGGNGTGCTTAACACATGCAAGTCGAACGATGAANCCTTTCGGGGTGGATTAGTGGCGAACGGGTGAGTAAC
ACGTGGGCAATCTGCCCTTCACTCTGGGACAAGCCCTGGAAACGGGGTCTAATACCGGATAACACTCTGTCCCGC
ATGGGACGGGGTTAAAAGCTCCGGCGGTGAAGGATGAGCCCCGCGCCTATCAGCTTGTTGGTGGGGTGATGGCCT
ACCAAGGCGACGACGGGTAGCCGGCCTGAGAGGGCGACCGGCCACACTGGGACTGAGACACGGCCAGACTCCTA
CGGGAGGCGAGCAGTGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAGCGACGCCGCTGAGGGATGACGGCC
TTCGGGTTGTAAACCTCTTTCAGCAGGGAAGAAGCGTGTGTCGGGAATTATTGGGCGTAAAGAGCTCGTANGCGGC
GTGCCAGCAGCCGCGTAATACGTAGGGCGCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGCTCGTANGCGGC
TTGTACGTCGGATGTGAAAGCCCCGGGGCTTAACCCGGGTCTGCATTCGATACGGGCTAACTAAAATGTGGTAGG
GGAGATCGGAATTCCTGGTGTANCGGTGAATGCGCAGATATCAAGAGGAACANCGGTGGCGAANGCGGATCTCT
GGGCCATTACTGACGCTGAGGAGCGAAAGCGTGGGAGCGAACAGGATTAGATACCCTGGTAGTCCACGCCGTAA
ACGTTGGGAAGTGGTGTGGCGACATTCCACGTCGTCGGTGCCGCGAGCTAACGCATTAAGTTCCCCGCTGGGG
AGTACGGCCGCAAGGCTAAAGTCAAAGGAATTGACGGGGGGCCCGCACAAAGCAGCGGAGCATGTGGCTTAATTCTG
ACGCAACGCGAAGAACCTTACCAAGGCTTGACATATACCGGAAAGCATCAGAGATGGTGCCCCCTTGTGGTTCGG
TATACAGGTGGTGCATGGCTGTCGTCAGCTCGTGTGAGATGTTGGGTAAAGTCCCGCAACGAGCGCAACCCCT
TGTTCTGTGTTGCCAGCATGCCCTTCGGGGTGATGGGACTCACAGGAGACTGCCGGGGTCAACTCGGAGGAAGG
TGGGGACGACGTCAAGTCATCATGCCCTTATGTCTTGGGCTGCACACGTGCTACAATGGCCGGTACAATGAGCT
GCGATGCCGCGAGGCGGAGCGAATCTCAAAAAGCCGGTCTCAGTTCCGATTGGGGTCTGCAACTCGACCCCATGA
AGTCGgAGTTGCTAgTAATCgCAgATCAgCATTGCTGCGGTGAATACgTTNCCGGGGCTtGTACAcACCGCCCGT
CACGTCACgAAAGTCGGTAACACCCGAAGNCCGGTGGTCCAACCCCTTGTGGGAGGGAGACTGTCGAAGGTGGGA
CTGGCGATTGG

EN43

CTCAGCGTCACTATCGGCCCAAAANACCGCCTTCGCCACCGGTGTTCTCCTGATATCTGCGCATTTACCCGCTA
CACCAGGAAATTCANTCTCCCCTACCGAACTCTANCTGCCCGTATCAACCGCAGGCTTGGGGTTAAGCCCCAA
TTTTTCACGGTCAACGCNACAAGCCGCTACAAGCTCTTTACGCCCAATAAATCCGGACAACGCTCGCACCCCTAC
TTCTTACCGCGGCTGCTGGCACTTATTTGGCCGGTGCTTCTTCTGCAGGTACCGTCACTCTCGCTTCGTCCCTGC
TNAAAAAGGTTTACAACCCGAAGGCCGTCATCCCTCACGCGGCNTCGCTGCATCAGGCTTCGGCCCATTTGTGCAA
TATTCCTCCACTGCTGCCTCCCGTAGGATTCTGGGCCGTNTCTCANTCCAGTGTGGCCGGTCGCCCTCTCAGGCC
GGCTACCCGTGCTGCCTTGGTAGGCCATCACCCACCAACAAGCTGATAGGCCGCGNAAGCCCATCCCAAGCCGA
AAAACCTTTCACCACCAGCCATGCGGCCAAAAATTCCTATTCCGGTATTAGCCCCCGTTTCNAAGGTTNTCCCAA
GCTTGGGGCAGGTTGCTCACTTTTTACTCACCCGTTCCCGCTCAATTACCCCNAAAGGGGNTTTCCTCAACTTGC
AT

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[FIGURE 19 CONTINUED]

EN47

CGCTGGCGGCGTGCTTAACACATGCAAGTCGAGCGGAAAGGCCCTTCGGGG₉TACTCGAGCGGGCGAACGGGTGAGT
AACACGTGAGTAATCTGCCCCAGGCTCTGGGATAGCCACCGGAAACGGTGATTAATACCGGATACGACAACCGAT
TGCATGATCTGGTTGTGGAAAGTTTTTCGGCCTGGGATGTGCTCGCGGCCCTATCAGCTTGTGGTGAGGTAATGG
CTCACCAAGGCTTCGACGGGTAGCCGGCCTGAGAGGGTGACCGGCCACACTGGGACTGAGACACGGCCCAGACTC
CTACGGGAGGCAGCAGTGGGGAATATTGGACAATGGGCGGAAGCCTGATCCAGCAACGCCCGCTGAGGGATGACG
GCCTTCGGGTTGTAAACCTCTTTCAGCACAGACGAAGCGCAAGTGACGGTATGTGCAGAAGAAGGACCGGCCAAC
TACGTGCCAGCAGCCGCGGTAATACGTAGGGTCCGAGCGTTGTCCGGAATTATTGGGCGTAAAGGGCTCGTAGGC
GGTCTGTGCGCTCGGGAGTGAAAACAGGTGCTTAACACCTGGCCTGCTTTCGATACGGGCAGACTAGAGGTACT
CAGGGGAGAATGGAATTCCT₉GTGTAGCGGTGAATGCGCAGATATTCAGGAGGAACACCGGTGGCGAAGGCGGT
TCTCTGGGAGTATCCTGACGCTTGAAGAGCGAAAGTGTGGGGAGCGAACAGGATTAGATANTNTGGTAGTCCACA
CCGTAAAC₉TTGGGCGCTAG₉TGTGGGACACATTCCACGTGTTCCGTGCCGCAGCTAACGCATTAAGCGCCCCGC
CTGGGAGTACGGCCGCAAGGCTAAACTCAAAGGAATTGACGGGGGCCCGCACAAAGCGGCGGAGCATGCGGATT
AATTCGATGCAACGCGAAGAACCTTACCTGGGTTTGACATACACCGGAAAGCCGTAGAGATACGGCCCCCTTTTAG
TCGGTGTACAGGTGGTGCATGGCTGTCGTGAGCTCGTGTGAGATGTTGGGTTAAGTCCCGAACGAGCGCAA
CCCTCGTCCTATGTTGCCAGCAATTCGGTTGGGACTCATAGGAGACTGCCGGGGTCAACTCGGAGGAAGGTGGG
GATGACGTCAAGTCATCATGCCCTTATGTCCAGGGCTTCACGCATGCTACAATGGCCGGTACAAAGGGCTCGCA
TCCCGTGAGGGTGAGCGAATCCCAAAAAGCCGGTCTCAGTTTCGGATTGGGGTCTGCAACTCGACCCCATGAAGTC
GGAGTCGCTAGTAATCGCAGATCAGCAACGCTGCGGTGAATACGTTCCCGGGCCTTGACACACCGCCCGTCACG
TCACGAAAGTCGGCAACACCCGAAGCCAGTGGCCCAACCCTTGTGGGGGGAGCTGTCGAAGGTGGGGCTGGCGAT
TG

EN59

GGGNATTAGTGGGGAACGGGTGAGTAAAANGTGGCCANTTTCCCTGNATTTTGGACANCCCCNGGAAANGGNTT
NTAAACNGGATANTGACCACCTTGGCATCCAAGTTTTNGAAACTTCCGGCGGTGCAGGATGAGCCNGCGGCNTA
TNAGCTTGTGGNGAGGTAATGGNTCACCAAGGGGANGACGGGTAGCCGGCCTGAGAGGGGACCNGCCACANTGGG
ANTGAGANACGGCCAGANTCCTACGGGAGGCAGCAGTGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAGG
ANNCCGCGTGAGGGANGACGGCCTTNGGGTTGTAAACNTTTTTNAGCAGGGAAGAAGCGAAAGTGACGGTACCTG
CAGAAGAAGCGCCGGCTAAATAAGTGCCAGCAGCCGCGGTAATAAGTAGGGNGCGAGCGTTGTNCGGAATTATTG
GGNGTAAAGAGTTTGTAGGCGGNTTGTNAAGTNGGTTGTGAAAGCCCCGGGNTTAACCCCGGGTTTGCAGTTGAT
ACGGGCAGGNTAGAGTTCGGTAGGGGAGATNGGAATTCCTGGTGTAGCGGTGAAATGCGCAGATATCAGGAGGAA
CACCGGTGGCGAAGGCGGATCTCTGGGCCATTACTGACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAG
ATACCTTGGTAGTCCACGCGTAAACGGTGGGAAGTACGGTGTGGCGACATTCCACGTCGTCGGTGGCGCAGCTA
ACGCATTAAGTTCCCCGCCTGGGGAGTACGGCCGCAAGGCTAAACTCAAAGGAATTGACGGGGGGCCCGCACAAAG
CAGCGGAGCATGTGGCTTAATTCGACGCAACGCGAAGAACCTTACCAAGGCTTGACATACACCGGAAAGCATCAG
AGATGGTGCCCCCTTGTGGTGGTGTACAGGTGGTGCATGGCTGTCTCAGCTCGTGTGAGATGTTGGGTT
AAGTCCCGCAACGAGCGCAACCCTTGTCTGTGTTGCCAGCATGCCCTTCGGGGTGTGGGGACTCACAGGAGAC
CGCCGGGGTCAACTCGGAGGAAGGTGGGGACGACGTCAAGTCATCATGCCCTTATGTCTTGGGCTGCACACGTG
CTACAATGGCCGGTACAAAGAGCTGCGATACCGTGAGGTGGAGCGAATCTCAAAAAGCCGGTCTCAGTTCGGATT
GGGGTCTGCAACTCGANCCCATGAANTCGGAGTTGCTAATTAATCGCAAAATCAAGCATTGCTGGCGGTGAATAC
GTTCCC

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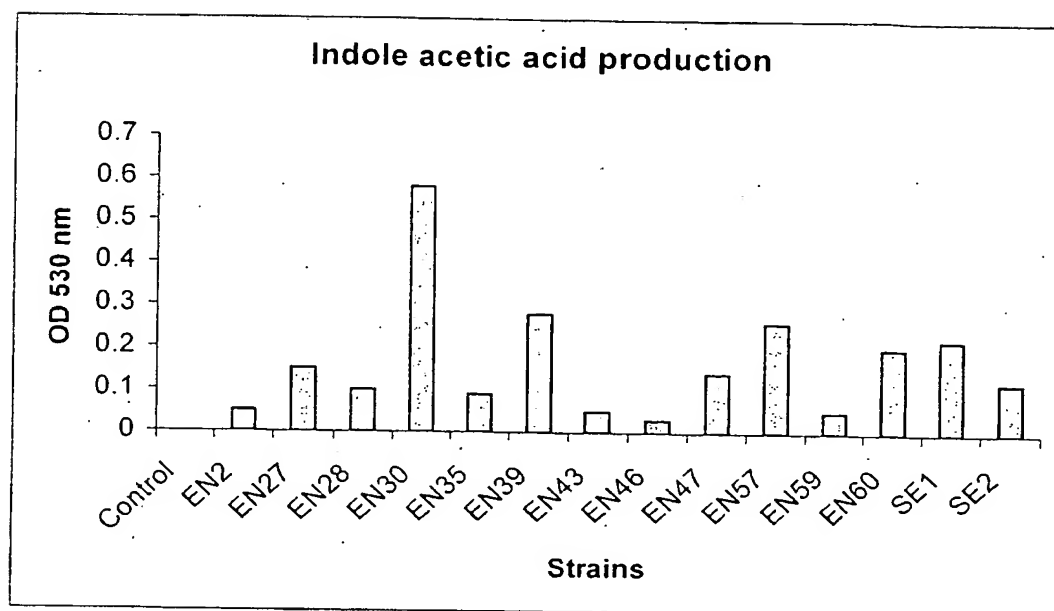
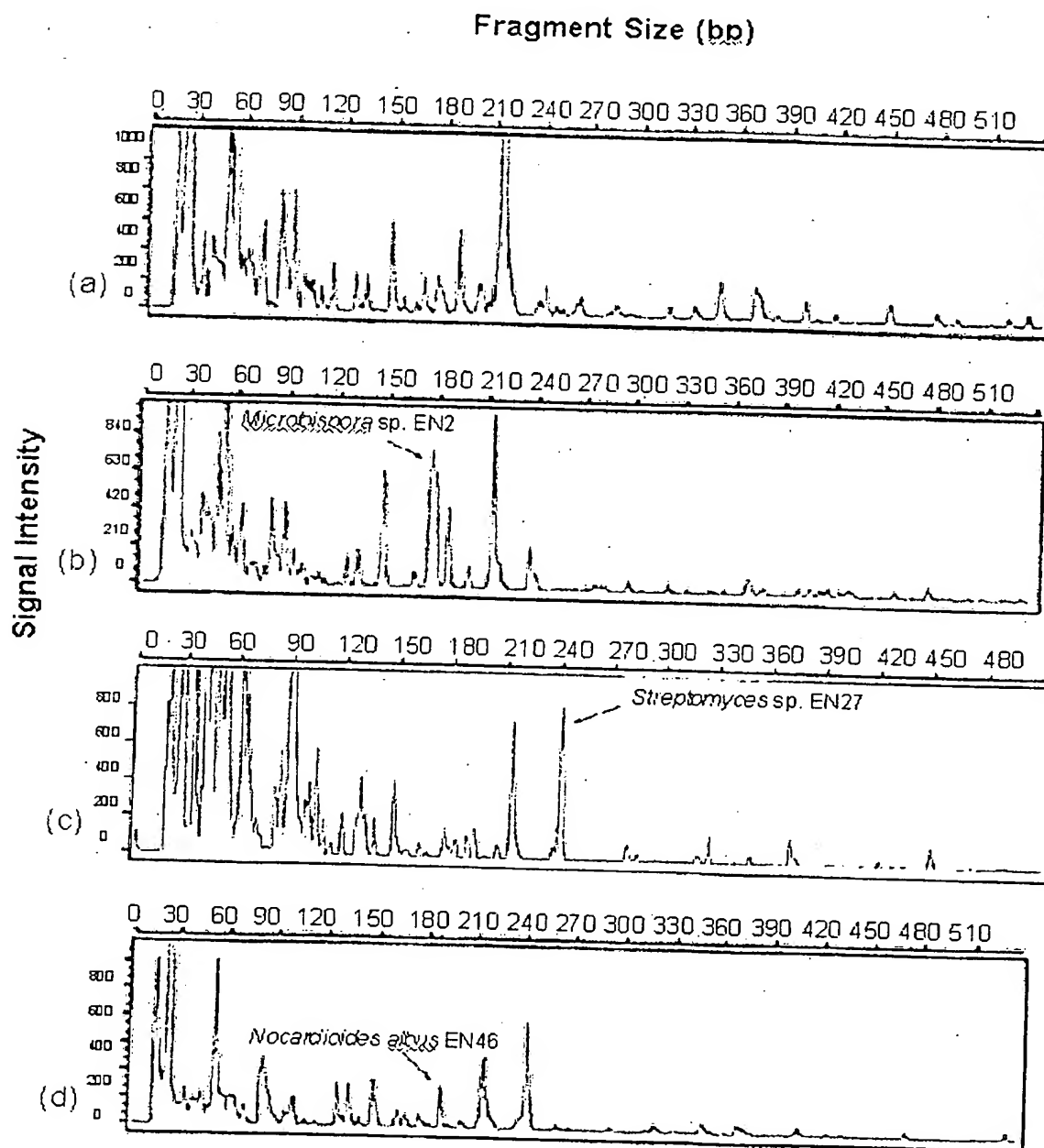


Figure 20

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Figure 21



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